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(72) Inventors: HIGGINS, Paul, J.; 8 Sheridan Aven ford, MA 02155 (US). POTTS, Barbara, J.; Street, Lexington, MA 02173 (US).	ue, Me 37 Gra	Published Mith international search report. Before the expiration of the time limit for amending the claims and to be republished in the event of the receipt of amendments.

(54) Title: HETEROCONJUGATE ANTIBODIES FOR TREATMENT OF HIV INFECTION

(57) Abstract

The invention features a heteroconjugate antibody which includes two binding functionalities. The first binding function is directed to a peripheral blood effector cell antigen, preferably CD-3. The second functionality is directed to a specific domain of an aids virus coat protein, preferably the V3 loop sequence of the gp120 envelope from HIV MN or a variant thereof. Methods of therapy are also presented where 20 ng/ml of a mixed culture of HIV infected cells (effector/CEM-ss cells) have a 80-90 % decrease in reverse transcriptase activity compared to an identical culture with a 3:1 ratio of effector cell:CEM-ss.

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HETEROCONJUGATE ANTIBODIES FOR TREATMENT OF HIV INFECTION Background of the Invention

This invention relates to the treatment of Human 5 Immunodeficiency Virus infection.

Human Immunodeficiency Virus (HIV), the etiologic agent of Acquired Immunodeficiency Syndrome (AIDS), is a retrovirus which infects certain immune system cells, including T4 lymphocytes and CD4⁺ cells of the monocyte/macrophage lineage. In the absence of effective treatment, the mortality rate for AIDS patients approaches 100% (Fauci, Science 239:617, 1988).

Well over 100 HIV variants have been identified. The amino acid sequence of the HIV envelope glycoprotein 15 gp120 is particularly variable; its amino acid sequence can vary by 20-25% from one strain to the next. addition to strain to strain variability, there is a more subtle variation in genome sequence caused by the high error rate of reverse transcriptase. The 20 misincorporation rate is high enough to introduce one error per genome per replication cycle. Consequently any particular viral isolate consists of a cohort of quasispecies. Further, the diversity and number of quasispecies apparently differs from one HIV variant to 25 another. There is substantial evidence that these quasispecies evolve in vivo. For example, successive viral isolates from an infected individual reveal substantial temporal fluctuations in the proportion of various quasispecies (Meyehans, Cell 58:901, 1989). There is also 30 evidence that neutralization-resistant HIV variants can

evidence that neutralization-resistant HIV variants can arise through single-base changes in the viral sequence encoding gp120 when HIV is grown in the presence of neutralizing antibodies (Reitz et al., Cell 54:57, 1988). Infected individuals initially mount a humoral and

cellular immune response against HIV, and there is reason to believe that an infected individual's immune response may actually encourage viral spread and the emergence of more resistant variants (McCune et al., Cell 64:351, 1991).

Human monoclonal antibodies directed against HIV proteins have been produced by hybridoma formation and Epstein Barr Virus transformation (Banapour et al., J. Immunol. 139:4027, 1987; Amadoci et al., AIDS Res. and 10 Human Retroviruses 5:73, 1989).

More recently cytotoxic hybrid proteins composed of a cytotoxin fused to part of the CD4 receptor have been proposed as a way to destroy cells expressing HIV encoded proteins. This approach relies on the fact that 15 the HIV envelope protein, gp120, recognizes the CD4 receptor, which is present on T4 lymphocytes and certain cells of the monocyte/macrophage lineage. Thus, a soluble derivative of CD4 might be used to target a cytotoxin to HIV infected cells that express surface 20 gp120. Chaudhary et al. (Nature 335:369, 1988) found that administration of a CD4-Pseudomonas exotoxin hybrid protein to a lymphocytic cell line chronically infected with HIV causes a decrease in overall protein synthesis. Till et al. (Science 242:1166, 1988) found that a CD4-25 ricin A fusion protein decreases DNA synthesis in cultures of chronically infected H9 cells. variation of this strategy, Capon et al. (Nature 337:529, 1989) designed a hybrid protein composed of soluble CD4 and the constant region of an antibody. This molecule is 30 designed to direct immune system response to gp120. Another molecule of this general type has been shown to activate complement (Traunecker et al. Nature 339:78, 1989).

Heteroconjugate molecules consisting of two
35 covalently joined antibodies or an antibody covalently

joined to a cell- or virus-targeted protein have been proposed as a means by which to target cytotoxic cells to undesirable cells such as tumor cells and virally infected cells. Segal et al. (U.S. Patent No. 4,676,980) 5 suggest the use of cross-linked hetero-antibodies to target immune system cells to unwanted or detrimental cells. Fanger et al. (PCT publication WO91/00360) have proposed such heteroconjugates for treatment of AIDS. particular, Fanger et al. suggest the use of a high 10 affinity Fc7 receptor-specific antibody fused to CD4 (or the CD4 binding domain of gp120) for AIDS therapy. Fanger et al. also suggest the use of heteroantibodies consisting of an high affinity Fc7 receptor-specific antibody fused to an HIV-specific antibody such as anti-15 gp120 antibody for AIDS therapy. Zarling et al. (EP Publication No. 03089.36) described heteroconjugate antibodies consisting of an antibody specific for an HIV antigen that is expressed on HIV infected cells crosslinked to a second antibody which is specific for an 20 effector cell of the peripheral blood and which is capable of killing HIV infected cells.

Summary of the Invention

In general, the invention features a heteroconjugate antibody which includes a first and a second portion joined together covalently, the first portion includes an antibody directed against an antigen present on the surface of an effector cell of the peripheral blood, the second antibody portion includes an antibody directed against a V3 loop sequence of the gp120 envelope protein of HIV MN or a HIV MN viral variant expressed on the surface of HIV-infected cells, wherein the heteroconjugate antibody at an initial concentration of 20 ng/ml in a first mixed cell culture which includes effector cells and CEM-ss cells infected with HIV-MN

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decreases the reverse transcriptase activity of the first mixed culture cell by at least 80% compared to the reverse transcriptase activity of an otherwise identical second mixed cell culture which includes effector cells and the CEM-ss cells infected with HTV-MN, wherein the effector cells are in 3-fold excess over the CEM-ss cells in the first and second mixed cell cultures, the reverse transcriptase activity is measured ten days after infection, the heteroconjugate antibody and the effector cells are added to the CEM-ss cells in the first mixed cell culture 18 hours after infection, and the first and second cell cultures are infected with 100-1000 infectious units of HIV-MN.

In a preferred embodiment, the decrease in the 15 reverse transcriptase activity of the first cell culture is greater than 90% compared to the reverse transcriptase activity of the second mixed cell culture.

In another preferred embodiment, the heteroconjugate antibody at an initial concentration of 20 200 ng/ml in a first mixed cell culture which includes effector cells and CEM-ss cells infected with an HIV strain other than HIV-MN decreases the reverse transcriptase activity of the first mixed culture cell by at least 50% compared to the reverse transcriptase 25 activity of an otherwise identical second mixed cell culture which includes effector cells and said CEM-ss cells infected with the HIV strain other than HIV-MN, wherein the effector cells are in 3-fold excess over said CEM-ss cells in the first and second mixed cell cultures, 30 the reverse transcriptase activity is measured ten days after infection, the heteroconjugate antibody and the effector cells are added to said CEM-ss cells in the first mixed cell culture 18 hours after infection, and the first and second cell cultures are infected with 100-

- 5 -

1000 infectious units of the HIV strain other than HIV-MN.

In yet another preferred embodiment, the heteroconjugate antibody binds to the V3 loop of an HIV 5 strain other than HIV-MN.

In other preferred embodiments, the effector cell is chosen from the group consisting of cytotoxic T lymphocytes, neutrophils, monocytes/macrophages, and large granular lymphocytes; and the antigen present on the surface of an effector cell is CD3.

In a another preferred embodiment, the heteroconjugate antibody at an initial concentration of 20 ng/ml in a first mixed cell culture which includes effector cells and CEM-ss cells infected with HIV-III, 15 decreases the reverse transcriptase activity of the first mixed culture cell by at least 80% compared to the reverse transcriptase activity of an otherwise identical second mixed cell culture which includes effector cells and the CEM-ss cells infected with HIV-III, wherein the 20 effector cells are in 3-fold excess over the CEM-ss cells in the first and second mixed cell cultures, the reverse transcriptase activity is measured ten days after infection, the heteroconjugate antibody and the effector cells are added to the CEM-ss cells in the first mixed 25 cell culture 18 hours after infection, and the first and second cell cultures are infected with 100-1000 infectious units of HIV-III,

In yet another preferred embodiment, the heteroconjugate antibody at an initial concentration of 20 ng/ml in three or more mixed cell cultures each of which includes effector cells and CEM-ss cells infected with one of the HIV strains: Alabama, Duke 6587-5, Duke 6587-7, Duke 7887-7, SF2, WMJ2, and IIIB, decreases the reverse transcriptase activity of each of the mixed cell culture by 80% compared to the reverse transcriptase

activity of an otherwise identical mixed cell culture which includes effector cells and CEM-ss cells infected with the same strain of HIV, wherein the effector cells are in 3-fold excess over the CEM-ss cells in the first and second mixed cell cultures, the reverse transcriptase activity is measured ten days after infection, the heteroconjugate antibody and the effector cells are added to the CEM-ss cells in the first mixed cell culture 18 hours after infection, and the first and second cell cultures are infected with 100-1000 infectious units of the strain of HIV.

In a related aspect, the invention features a heteroconjugate antibody which includes a first and a second portion joined together covalently, the first 15 portion includes an antibody directed against an antigen present on the surface of an effector cell of the peripheral blood, the second antibody portion includes an antibody directed against a V3 loop sequence of the gp120 envelope protein of HIV MN or a HIV MN viral variant 20 expressed on the surface of HIV-infected cells, wherein the heteroconjugate antibody at an initial concentration of 10 ng/ml in a first mixed cell culture which includes the effector cells and CEM-ss cells infected with HIV-MN decreases the reverse transcriptase activity of the first 25 mixed culture cell by at least 80% compared to the reverse transcriptase activity of an otherwise identical second mixed cell culture which includes the effector cells and the CEM-ss cells infected with HIV-MN, wherein the effector cells are in 3-fold excess over the CEM-ss 30 cells in the first and second mixed cell cultures, the reverse transcriptase activity is measured ten days after infection, the heteroconjugate antibody and the effector cells are added to the CEM-ss cells in the first mixed cell culture 18 hours after infection, and the first and

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second cell cultures are infected with 100-1000 infectious units of HIV-MN.

In a related aspect, the invention feature a heteroconjugate antibody which includes a first and a 5 second portion joined together covalently, the first portion includes an antibody directed against an antigen present on the surface of an effector cell of the peripheral blood, the second antibody portion includes an antibody directed against a V3 loop sequence of the gp120 10 envelope protein of HIV MN or a HIV MN viral variant expressed on the surface of HIV-infected cells, wherein the heteroconjugate antibody at an initial concentration of 5 ng/ml in a first mixed cell culture which includes effector cells and CEM-ss cells infected with HIV-MN 16 decreases the reverse transcriptase activity of the first mixed culture cell by at least 80% compared to the reverse transcriptase activity of an otherwise identical second mixed cell culture which includes effector cells and the CEM-ss cells infected with HIV-MN, wherein the 20 effector cells are in 3-fold excess over the CEM-ss cells in the first and second mixed cell cultures, the reverse transcriptase activity is measured ten days after infection, the heteroconjugate antibody and the effector cells are added to the CEM-ss cells in the first mixed 25 cell culture 18 hours after infection, and the first and second cell cultures are infected with 100-1000 infectious units of HIV-MN.

In another related aspect, the invention features a heteroconjugate antibody which includes a first and a second portion joined together covalently, the first portion includes an antibody directed against an antigen present on the surface of an effector cell of the peripheral blood, the second antibody portion includes an antibody directed against a V3 loop sequence of the gp120 envelope protein of HIV MN or a HIV MN viral variant

expressed on the surface of HIV-infected cells, wherein the heteroconjugate antibody at an initial concentration of 1 ng/ml in a first mixed cell culture which includes effector cells and CEM-ss cells infected with HIV-MN 5 decreases the reverse transcriptase activity of the first mixed culture cell by at least 80% compared to the reverse transcriptase activity of an otherwise identical second mixed cell culture which includes effector cells and the CEM-ss cells infected with HIV-MN, wherein the 10 effector cells are in 3-fold excess over the CEM-ss cells in the first and second mixed cell cultures, the reverse transcriptase activity is measured ten days after infection, the heteroconjugate antibody and the effector cells are added to the CEM-ss cells in the first mixed 15 cell culture 18 hours after infection, and the first and second cell cultures are infected with 100-1000 infectious units of HIV-MN.

In a related aspect, the invention features a heteroconjugate antibody which includes a first and a second portion joined together covalently, the first portion includes an antibody directed against an antigen present on the surface of an effector cell of the peripheral blood, the second antibody portion includes an antibody directed against the amino acid sequence GPGRAF.

In a preferred embodiment, the heteroconjugate antibody at an initial concentration of 20 ng/ml in a first mixed cell culture which includes effector cells and CEM-ss cells infected with HIV-MN decreases the reverse transcriptase activity of the first mixed culture cell by at least 80% compared to the reverse transcriptase activity of an otherwise identical second mixed cell culture which includes cells and the CEM-ss cells infected with HIV-MN, wherein the effector cells are in 3-fold excess over the CEM-ss cells in the first and second mixed cell cultures, the reverse transcriptase

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activity is measured ten days after infection, the heteroconjugate antibody and the effector cells are added to the CEM-ss cells in the first mixed cell culture 18 hours after infection, and the first and second cell cultures are infected with 100-1000 infectious units of HIV-MN.

In a related aspect, the invention features a heteroconjugate antibody which includes a first and a second portion joined together covalently, the first 10 portion includes an antibody directed against an antigen present on the surface of an effector cell of the peripheral blood, the second antibody portion includes an antibody directed against the amino acid sequence IXIGPGR, wherein X = any amino acid.

In a preferred embodiment, the heteroconjugate 15 antibody at an initial concentration of 20 ng/ml in a first mixed cell culture which includes effector cells and CEM-ss cells infected with HIV-MN decreases the reverse transcriptase activity of the first mixed culture 20 cell by at least 80% compared to the reverse transcriptase activity of an otherwise identical second mixed cell culture which includes effector cells and the CEM-ss cells infected with HIV-MN, wherein the effector cells are in 3-fold excess over the CEM-ss cells in the 25 first and second mixed cell cultures, the reverse transcriptase activity is measured ten days after infection, the heteroconjugate antibody and the effector cells are added to the CEM-ss cells in the first mixed cell culture 18 hours after infection, and the first and 30 second cell cultures are infected with 100-1000 infectious units of HIV-MN.

In a related aspect, the invention features a heteroconjugate antibody which includes a first and a second portion joined together covalently, the first portion includes an antibody directed against an antigen

present on the surface of an effector cell of the peripheral blood, the second antibody portion includes an antibody directed against the amino acid sequence QARILAVERYLKDQQLLGIWGCSGKLIC.

In a preferred embodiment, the heteroconjugate 5 antibody at an initial concentration of 20 ng/ml in a first mixed cell culture which includes effector cells and CEM-ss cells infected with HIV-MN decreases the reverse transcriptase activity of the first mixed culture 10 cell by at least 80% compared to the reverse transcriptase activity of an otherwise identical second mixed cell culture which includes effector cells and the CEM-ss cells infected with HIV-MN, wherein the effector cells are in 3-fold excess over the CEM-ss cells in the 15 first and second mixed cell cultures, the reverse transcriptase activity is measured ten days after infection, the heteroconjugate antibody and the effector cells are added to the CEM-ss cells in the first mixed cell culture 18 hours after infection, and the first and 20 second cell cultures are infected with 100-1000 infectious units of HIV-MN.

In other preferred embodiments, the effector cell is chosen from the group consisting of cytotoxic T lymphocytes, neutrophils, monocytes/macrophages, and large granular lymphocytes; and the antigen present on the surface of an effector cell is CD3.

In another aspect, the invention features a pharmaceutically acceptable composition which includes a pharmaceutically effective amount of a heteroconjugate 30 antibody described above.

In a related aspect, the invention features a method for treating a patient infected with HIV, the method includes administering to the patient the above-described pharmaceutically acceptable composition.

In another aspect, the invention features an HIV-targeted effector cell which includes: (a) an effector cell expressing a cell surface antigen; and (b) an above-described heteroconjugate antibody.

In a related aspect, the invention features a method for treating a patient infected with HIV; the method includes administering to the patient the above-described HIV-targeted effector cell.

The MN prototype virus is defined by a particular amino acid subsequence within the V3 loop region of the gp120 envelope protein having positions A₁-A₁₇:K-R-K-R-I-H-I-G-P-G-R-A-F-Y-T-T-K. (Amino acid sequences are presented in the standard single-letter code throughout.) MN viral variants are variant which exhibit complete amino acid sequence homology at residues I-G-P-G-R, i.e., at positions A₇ through A₁₁, and at least 36% homology with the remaining 12 amino acids of the HIV-MN sequence given above.

By "directed against" is meant that an antibody binds to 20 the indicated antigen. The V3 loop of gp120 is defined as the 36 amino acid region from amino acid 303 to 338, inclusive, according to the gp120 numbering scheme of Ratner et al. (Nature 313:277, 1985).

The heteroconjugate antibodies of the invention
25 are highly effective; even at low concentrations they are
capable of nearly eliminating viral replication as judged
by a reverse transcriptase assay. The preferred
heteroconjugate antibodies are those which are effective
against more than one strain.

Other features and advantages of the invention will be apparent from the following description of the preferred embodiments thereof, and from the claims.

30

<u>Detailed Description</u>

The drawings are first briefly described.

Figure 1 is a graphical representation of the effect of a mixture of unconjugated OKT3 antibody and 59.1 antibody (filled diamonds) and OKT3/59.1 heteroconjugate antibody (open squares) on the reverse transcriptase activity of CEM-ss cells infected with HIV-III_B in the presence of cytotoxic T-lymphocytes. Reverse transcriptase activity (cpm/10 μ l) is presented as a function of the initial antibody concentration (ng/ml) in the cell culture.

10 Figure 2 is a graphical representation of the effect of a mixture of unconjugated OKT3 antibody and 59.1 antibody (filled diamonds) and OKT3/59.1 heteroconjugate antibody (open squares) on the reverse transcriptase activity of CEM-ss cells infected with HIV-15 MN in the presence of cytotoxic T-lymphocytes. Reverse transcriptase activity (cpm/10 μ l) is presented as a function of the initial antibody concentration (ng/ml) in the cell culture.

effect of a mixture of unconjugated OKT3 antibody and 59.1 antibody (open circles) and OKT3/59.1 heteroconjugate antibody (filled circles) on the reverse transcriptase activity of CEM-ss cells infected with HIV-III_B in the absence of cytotoxic T-lymphocytes. The reverse transcriptase activity of HIV-III_B infected CEM-ss cells in the presence of cytotoxic lymphocytes only (filled triangle); HIV-III_B infected CEM-ss cells alone (open triangle); and uninfected CEM-ss cells alone (filled square) is also indicated. Reverse transcriptase activity (cpm/10 µl) is presented as a function of the initial antibody concentration (ng/ml) in the cell culture (except for those cases in which no antibody was added).

Figure 4 is a graphical representation of the 35 effect of the ratio of cytotoxic T-lymphocytes to HIV-MN

infected CEM-ss cells on reverse transcriptase activity in the presence of 1 μ g/ml OKT3/59.1 heteroconjugate antibody. Reverse transcriptase activity (cpm/10 μ l) is presented as a function of the cytotoxic lymphocyte to 5 CEM-ss cells (log₁₀ scale).

Figure 5 is a set of graphs which illustrate the effect of cytotoxic T-lymphocytes and OKT3/59.1 heteroconjugate antibody (open squares) and a mixture of unconjugated OKT3 antibody and 59.1 antibody (filled 10 triangles) on the reverse transcriptase activity of CEM-ss cells infected with HIV-MN (panel A), HIV-Alabama (panel B), HIV-Duke 7887-7 (panel C), HIV-Duke 6587-5 (panel D), HIV-Duke 6587-7 (panel E), HIV-III_B (panel F), HIV-SF2 (panel G), and HIV-WMJ2 (panel H). In each case, 15 a control is included where neither antibodies or cytotoxic T-lymphocytes were added (filled circles). Reverse transcriptase activity (cpm/10 µl) is presented as a function of the number of days post-infection.

Figure 6 is a graphical representation of the
20 effect of a mixture of conjugated OKT3 antibody and 6C5
antibody (open circles) and OKT3/6C5 heteroconjugate
antibody (filled circles) on the reverse transcriptase
activity of HIV-III_B infected CEM-ss cells. Reverse
transcriptase activity (cpm/10 µl) is presented as a
25 function of the initial antibody concentration (ng/ml) in
the cell culture.

Heteroconjugate Antibodies for AIDS Therapy

The molecules of the invention are heteroconjugate antibodies produced by covalently attaching a first

30 antibody which is directed against an antigen present on the surface of a cytotoxic immune effector cell capable of killing an HIV infected cell to a second antibody which is directed against an HIV antigen present on the surface of HIV infected cells.

The heteroconjugate antibodies of the invention are highly potent. Even at relatively low concentrations, these heteroconjugate antibodies are capable of substantially reducing HIV activity in a mixed cell culture of HIV infected cells and effector cells. The most preferred heteroconjugate antibodies are those which are both highly potent and broadly reactive. Broadly reactive heteroconjugate antibodies are those which are effective against more than one strain of HIV.

To example, a broadly reactive heteroconjugate antibody might be effective against HIV-MN and HIV-SF2 or HIV-MN and HIV-WMJ2, or HIV-MN and HIV-III_B.

The portion of the heteroconjugate antibody which is directed against a cytotoxic immune effector cell

15 capable of killing HIV infected cells recognizes an antigen present on the surface of cells such as: cytotoxic T-lymphocytes, monocytes/macrophages, large granular lymphocytes (including cells and NK cells), and neutrophils. Preferably, the immune effector cell
20 directed antibody binds to an antigen on the surface of the effector cell in a manner which triggers cytolytic activity. For example, the antigen recognized can be the CD3 receptor or the CD16 (Fc) receptor. Less preferred are antibodies directed to receptors which require

25 multiple signals to initiate cytolytic activity (e.g., the CD2 and CD28 receptors).

The portion of the heteroconjugate antibody which is directed against an antigen present on the surface of HIV infected cells preferably recognizes: (1) an epitope within the V3 loop sequence of the gp120 envelope protein of the MN prototype of HIV-1 (HIV-MN); (2) an epitope within the V3 loop sequence of the gp120 envelope protein of a viral variant of the MN prototype of HIV-1; or (3) an epitope within the portion of gp41 between amino acids 584 to 611, inclusive.

The V3 loop of gp120 is the 36 amino acid region from amino acid 303 to 338, inclusive, according to the gp120 numbering scheme of Ratner et al. (Nature 313:277, 1985). The MN prototype of HIV-1 is defined by the following amino acid subsequence within the V3 loop of gp120: K-R-K-R-I-H-I-G-P-G-R-A-F-Y-T-T-K (A¹-A¹⁷). MN viral variants are variants which exhibit complete amino acid homology at residues I-G-P-G-R, i.e., positions A⁷ through A¹¹, and at least 36% homology with the remaining 10 12 amino acids of the MN sequence given above.

The above-described HIV-directed antibodies are good candidates for use in generating heteroconjugate antibodies which are highly potent. In some cases, however, the heteroconjugates formed will not be highly 15 potent. Ultimately the usefulness of a particular HIVdirected antibody for production of a heteroconjugate antibody of the invention can only be assessed by producing a heteroconjugate antibody, for example by covalently linking the HIV-directed antibody to an anti-20 CD3 antibody, and measuring the potency of the heteroconjugate antibody in an appropriate assay. particular HIV-directed antibody has been shown to be useful for generating a potent heteroconjugate antibody, it can be used to generate other heteroconjugate 25 antibodies by covalently linking it to other effector cell antigen-directed antibodies.

As discussed above, the most preferred heteroconjugate antibodies are those which are broadly reactive as well as highly potent. HIV-directed

30 antibodies which recognize: (1) an epitope having the sequence G-P-G-R-A-F; (2) an epitope having the sequence I-X-I-G-P-G-R, where X is any amino acid; or (3) an epitope within the portion of gp41 between amino acids 584 to 611 (according to the numbering scheme of Ratner et al., supra), inclusive are likely to be useful for

generating highly potent and broadly reactive heteroconjugate antibodies. This does not imply that antibodies recognizing other epitopes within the V3 loop of HIV-MN, the V3 loop of an HIV-MN viral variant, or gp41 cannot be used to produce a highly potent, broadly reactive heteroconjugate antibodies.

Described below are techniques for generating and screening HIV-directed antibodies useful for preparation of heteroconjugate antibodies, methods for the 10 preparation of heteroconjugate antibodies, and methods for assessing the potency and breadth of reactivity of heteroconjugate antibodies. In order to generate broadly reactive heteroconjugate antibodies it is useful to select HIV-directed antibodies which recognize a broad 15 range of HIV strains (i.e., antibodies which are not strain specific). It is also useful to select antibodies which are directed against the amino acid sequence: G-P-G-R-A-F; or the amino acid sequence: I-X-I-G-P-G-R, where X is any amino acid; or an epitope within the portion of 20 gp41 from amino acid 584 to amino acid 611. These antibodies can be identified using standard epitope mapping techniques as described below.

useful HIV-directed antibodies include: (1) generation of hybridomas and selection of hybridomas producing reactive antibodies; (2) selection of hybridomas producing antibodies capable of binding to cells expressing HIV envelope protein; (3) amplification and purification of selected monoclonal antibodies; (4) analysis of antibody reactivity using gp120 V3 loop peptides or gp41 derived peptides; and (5) epitope mapping. As mentioned above not all of these steps are essential. It is possible to simply follow steps 1 through 3 and use the purified antibodies to prepare heteroconjugate antibodies whose potency and breadth of reactivity can be analyzed using

the reverse transcriptase assay described herein below.

To generate a heteroconjugate antibody of the invention,
a purified HIV-directed antibody is covalently attached
to an antibody directed against an immune effector cell.

The potency and reactivity of heteroconjugate antibodies
can be measured in a mixed cell culture of effector cells
and HIV-infected cells using a reverse transcriptase
assav.

Heteroconjugate antibodies may be formed using any convenient cross-linking method. Suitable cross-linking methods include: SPDP, SPDP and SMCC, and biotin-avidin. Segal et al. (U.S. Patent 4,676,980) describes a number of cross-linking techniques. Alternatively, the antibodies can be linked by the generation of bispecific antibodies via hybrid-hybridomas (Suresh et al., Methods in Enzymology 121:210, 1896) or by genetic engineering. Preparation of HIV-directed Antibodies

Antibodies useful for the preparation of the heteroconjugate molecules of the invention may be generated and screened as described below. Methods for preparing and analyzing antibodies directed towards the V3 loop of HIV-MN of an HIV-MN viral variant are also described in U.S. Application No. 07/665,306, filed March 6, 1991, hereby incorporated by reference.

25 Preparation of the Immunogen

One group of HIV-directed antibodies useful for production of heteroconjugate antibodies recognize sequences within the V3 loop of HIV-MN or an HIV-MN viral variant. Accordingly, the immunogen used to generate these antibodies can include: gp160, gp120, fragments of gp120 or gp160 which include all or part of the V3 loop, or synthetic peptides which include all or part of the V3 loop. In all cases the V3 loop sequences is that of HIV-MN or an HIV-MN viral variant. Preferred immunogens for

generating V3 loop directed antibodies include the RP70 peptide formed into a closed loop (described below).

The other group of HIV-directed antibodies useful for production of heteroconjugate antibodies recognize sequences within the region of gp41 spanning amino acids 584-611. The immunogens used to generate these antibodies can include: gp160, gp41, and fragments of gp160 or gp41 which include all or part of the sequence lying between amino acids 584 and 611 of gp41, i.e., Q-10 A-R-I-L-A-V-E-R-Y-L-K-D-Q-Q-L-L-G-I-W-G-C-S-G-K-L-I-C.

The immunizing peptide, polypeptide or protein may be in linear form or alternatively may contain the V3 loop formed into a closed loop by creation of a disulfide bond between cysteine residues at the termini of the V3 loop sequence. If the immunizing peptide contains more than one V3 loop, each may be separately formed into a loop through disulfide bonding.

sequences can be synthesized by automated peptide
sequences can be synthesized by automated peptide
synthesis using an automated peptide synthesizer. Intact
recombinant gp160 envelope polypeptide can be produced in
insect cells using a baculovirus expression system and
purified as described in Rusche et al., U.S. Application
No. 091,481, filed August 31, 1987, assigned to the same
assignee as the present invention, hereby incorporated by
reference.

synthetic peptides or protein fragments to be used as immunogens can be either unconjugated or conjugated to an immunogenic carrier, e.g., keyhole limpet hemocyanin (KLH) or ovalbumin, using succinyl maleimidomethyl cyclohexanylcarboxylate (SMCC) as a conjugation agent (Yoshitake et al., J. Biochem. 92:1413, 1982), as follows.

Briefly, 1 mg of SMCC dissolved in 50 μ l of 35 dimethylformamide is added to 6 mg of carrier (at a

30 RP339 (RF)

concentration of 10-20 mg/ml in 0.1M NaPO₄, pH 6.5) and incubated at room temperature for 0.5 h. The solution is then passed through a Sephadex G-25 column to remove excess unreacted SMCC and 2 mg of peptide is added

5 (suspended in a degassed solution of 0.1M NaPO₄, pH 8, 1mM EDTA at a concentration of 10 mg/ml). The solution is mixed by N₂ gas and incubated at 4°C overnight. The sample is then dialyzed in 6M urea, 0.1M NaPO₄, pH 7 until the precipitate dissolves. The sample is next eluted through a BioGel P-10 column equilibrated in 6M urea, 0.1M NaPO₄. The voided protein is collected and dialyzed in distilled H₂O.

The sequences of several peptides (RP142, RP70, RP342, RP100, RP102, RP108, RP123c, and RP174c) useful in immunogens are shown in Table 1. This list is not meant to be exhaustive; it merely lists a few of the peptides which may be used as immunogens.

Table 1: Examples of Peptides Useful as Immunogens YNKRKRIHIGPGRAFYTTKNIIG (C) RP142 20 RP342 IHIGPGRAFYT RP70 INCTRPNYNKRKRIHIGPGRAFYTTKN I I G T I R Q A H C N I S (SGG) TRKGIHIGPGRAIY (GGSC) RP100 RP102 (SGG) TRKSISIGPGRAF (GGSC) . 25 RP108 (SGG) HIGPGRAFYATG (GGSC) RP123c (C) HIGPGRAF (C) RP135 (III_B) NNTRKSIRIQRGPGRAFVTIGKIG (C) (C) NNTRKSIRIQRGPGRAFVTIGKIG RP174c (C)

Note: Amino acids in parentheses are not in the natural sequence of the indicated isolate

ITKGPGRVIY (C)

Peptides RP70, RP123c, and RP174c can be formed into closed loops by creation of a disulfide bond between the two cysteine residues near the ends of the amino acid sequence. A method for creating such a bond is described in Zhang et al. (Biochemistry 27:3785, 1988).

The peptides were prepared for immunization by emulsification in complete Freund's adjuvant according to standard techniques. (CFA, Difco Labs, Grand Island, NY). Generation of HIV-Directed Antibodies

HIV-directed antibodies were prepared by 5 intraperitoneal immunization of mouse strains (Balb/c, C57BL/6, A.SW, B10.BR, or B10.A, Jackson Labs., Bar Harbor, ME) with 10-50 μ g per mouse of circularized RP70 (Table 1) or recombinant gp160. The mice were given 10 booster immunizations of the immunogen, either in an emulsification of incomplete Freund's adjuvant or in soluble form, two to three times at two to four week intervals following the initial immunization. Mice were bled and the sera assayed for the presence of antibodies 15 reactive with the immunogen. Mice showing a strong serological response were boosted and, 3-5 days later, spleen cells from these mice were fused with NS-1 (A.T.C.C. No. TIB18), SP2-0 (A.T.C.C. No. CRL8287, CRL8006), or P3.X63.AG8.653 myeloma cells incapable of 20 secreting both heavy and light immunoglobulin chains (Kearney et al., J. Immunol. 123:1548, 1979) by standard procedures based on the method of Kohler and Milstein, (Nature 256:495, 1975).

Supernatants from hybridomas which appeared 6-21
25 days after fusion were screened for production of
antibodies by an ELISA screening assay, as follows. The
RP70 peptide was used to screen RP70 generated
hybridomas, and a peptide whose amino acid sequence is
identical to that of residues 567-647 of gp41, was used
30 to screen gp160 generated hybridomas.

Each well of a 96-well Costar flat-bottom microtiter plate was coated with the peptide by placing a 50 μ l aliquot of a PBS solution containing the peptide at a final concentration of 0.1-10 μ g/ml in each well. The peptide solution was aspirated and replaced with PBS +

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0.5% BSA. Following incubation, the wells were aspirated, washed, and 50 μl of hybridoma supernatant was added. Following incubation, the wells were washed 3 times with PBS, and then incubated with 50 μl of an
5 appropriate dilution of goat anti-mouse immunoglobulin conjugated with horseradish peroxidase (HRP, Zymed Laboratories, San Francisco, CA). The wells were washed again 3 times with PBS and 50 μl of 1mM ABTS (2,2 azinobis (3-ethylbenzthiazoline-6-sulfonic acid) in 0.1M Na-10 Citrate, pH 4.2, to which a 1:1000 dilution of 30% H₂O₂ had been added), the substrate for HRP, was added to detect bound antibody. HRP activity was monitored by measuring the absorbance at 410nm.

Hybridomas that test positive by the ELISA method 15 can be tested for their ability to bind to cells which express the HIV envelope protein. In one such assay recombinant vaccinia virus expressing a the env gene of a particular HIV strain are used to infect cells of the CD4+ human T-lymphoma line, CEM-ss (AIDS Research and 20 Reference Reagent Program, Rockville, MD, catalog #776). Hybridoma supernatant (or purified antibodies) are incubated with the infected cells, and antibody binding is detected by indirect immune florescence using a secondary antibody and a florescence activated cell 25 sorter. As a control, binding to otherwise identical cells which do not express an HIV env gene is measured. Hybridomas producing antibodies which bind to env expressing cells (but not to non-expressing cells) are then selected for further characterization. Cells 30 expressing the env gene of any HIV strain may be prepared as described below.

In some cases (for neutralizing antibodies) an assay for inhibition of syncytia formation can be used to assess HIV-directed antibodies. In this assay the antibody is added to a mixture of HIV-infected and

uninfected cells and giant cell formation is monitored. This assay is described in detail in U.S. Application No. 07/665,306, filed March 6, 1991, hereby incorporated by reference.

5 <u>Preparation of HIV env Expressing Cells Using a</u> Recombinant Vaccinia Virus

An assay for binding to cells expressing an HIV env gene can employ cells infected with a vaccinia virus expressing an HIV env gene rather than actual HIV

10 infected cells. Construction of a recombinant vaccinia virus capable of expressing the full-length HIV envelope gene from a vaccinia virus promoter is described in EP Publication No. 0 243 029, hereby incorporated by reference. The recombinant vector pSC25, containing the

15 HIV env gene and the lacZ gene of E. coli expressed from a second vaccinia virus promoter, and flanked by vaccinia viral sequences which together encode thymidine kinase (TK), was used to produce the recombinant virus.

A recombinant vector that contains DNA encoding an envelope gene having the specificity of the HIV-MN variant was prepared by removing a 570 bp BglII fragment (encoding 180 amino acids) from the HIV-III_B env gene which spans the region of the VS loop in pSC25, and replacing it with the analogous BglII fragment from the HIV-MN env gene. The resulting plasmid, pSCR2502, contained a hybrid envelope gene which encoded an envelope protein having the principal neutralizing domain of the MN virus and the remainder of the env gene sequence from the HIV-III_B envelope.

A smaller region of the HIV-MN gp160 protein can be used in place of the 180 amino acid replacement just described; e.g., DNA encoding the 36 amino acid V3 loop from any HIV strain can be inserted into the envelope-encoding DNA in place of the corresponding III_B DNA sequence. Alternatively, a recombinant could be used

which contains the complete HIV-MN env gene. Multiple HIV envelope expressing strains are useful for assessing the specificity of an antibody.

The recombinant vector pSCR2502 was transfected into CV-1 host cells that had been pre-infected with vaccinia virus containing an intact TK gene. The HIV envelope gene was integrated into the viral DNA by homologous recombination between the TK sequences on the vector and the TK sequences within the viral genome.

- 10 Recombinants containing the HIV envelope gene were selected by infection of TK- cells and plating on media containing bromodeoxyuridine (BUdR) and X-gal. BUdR is toxic to TK+ cells and thus selects for TK- recombinants; X-gal is a chromogenic substrate cleaved by the product
- of the lacZ gene which results in the production of blue plaques where the lacZ gene is expressed and further identifies the recombinant virus which also contains the HIV-env gene.

Antibody Purification and Amplification

- 20 Hybridomas that tested positive for peptide binding in the ELISA assay were subcloned by the limiting dilution method. Hybridoma cells and irradiated splenocytes from nonimmunized syngeneic mice (final concentration 5 cells/ml and 2.5 x 10⁶ cells/ml,
- 25 respectively) were mixed and 200 μl of the mixed suspension were plated in microtiter wells to give 1 hybridoma cell per well. Subclones which appeared 7-14 days later were assayed again by the ELISA procedure described above. Representative positive subclones were 30 subcloned a second time.

The isotypes of the antibodies were determined by the ELISA method using goat anti-mouse-HRP preparations which corresponded to each of the five major mouse immunoglobulin isotypes (IgM, IgG1, IgG2A, IgG2B and 35 IgG3).

30

Purified antibodies were prepared by injecting hybridoma subclones that repeatedly tested positive by ELISA and syncytium inhibition assays intraperitoneally into pristane-primed syngeneic mice. The ascites which developed were recovered two to three weeks after injection and the monoclonal antibodies were purified as follows, using procedures which were dependent on the isotype of the antibody. Following elution, all IgG antibodies were dialyzed against PBS.

10 IgM antibodies were purified by 50% NH₂SO₄
precipitation of ascites fluid from mice injected with
the corresponding hybridoma cells, and then dialysis of
the precipitate against 4X PBS. The dialyzed antibody
was then passed over an Ultrogel A-6 column
15 (Biotechnics, Villeneuve-La-Garenne, France) preequilibrated with 4X PBS. The antibody-containing
fraction was identified using ELISA.

Ascites fluid containing IgG1 antibodies was diluted 4-fold in 0.1M Tris-HCl, 3M NaCl, pH 8.9, and 20 isolated by passage through a Protein A-Sepharose affinity column equilibrated with the same Tris-NaCl buffer. The antibody was eluted using 0.1M Na-Citrate, pH 6.0.

Ascites fluid containing IgG2 antibodies was
25 diluted two-fold in PBS, and then bound to a Protein-Asepharose affinity column equilibrated with PBS. It was
then eluted from the column with 0.15M NaCl, 0.1M acetic
acid, pH 3.0. Following elution, the antibody was
immediately neutralized by the addition of 1M Na₂HCO.

Ascites fluid containing IgG3 antibodies was diluted 4-fold in 0.1M Tris-HCl, 3M NaCl, pH 8.9, passed over a Protein-A-Sepharose affinity column, and antibody was eluted from the Protein A column with 0.15M NaCl, 0.1M acetic acid.

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Alternatively, all IgG subclasses can be purified by the following procedure. Ascites fluid is diluted 2-fold in 0.1M Tris-HCl, 3M NaCl pH 8.9, passed over Protein A Sepharose affinity column, and eluted with 5 0.15M NaCl, 0.1M acetic acid, pH 3.0.

Determination of Antibody Specificity Assays described below can be used for determination of the strain specificity of HIV-directed antibodies and to map the epitope recognized by HIV-10 directed antibodies. Some or all of these assays may be used to select HIV-directed antibodies for production of heteroconjugate antibodies. The assay for binding to cells expressing an HIV env gene described above can also be used to assess antibody specificity. The epitope 15 recognized by the V3-directed antibodies can be mapped using standard ELISA assays and competitive ELISA assays as described below. Peptides which are useful for ELISA assays include: (1) a series of 24 or 25-mers representing the V3 loop sequences from a variety of HIV 20 variants (Table 2); and (2) the MN substitution series, which includes a series of 12-mers corresponding to the MN V3 loop tip sequence (C)-K-R-I-H-I-G-P-G-R-A-F-Y-T-T-(C), each having an alanine residue substituted for one of the amino acids starting at the first arginine (R) 25 residue and proceeding to the tyrosine (Y) residue. the substitution series a glycine was substituted for the naturally occurring alanine. Antibody recognition of an epitope contained within the MN sequence is revealed by loss of binding of the antibody to an alanine-substituted

Competitive ELISA assays were performed as for standard ELISA assays with the following modifications. Prior to applying the antibody to the plate, the antibody preparation is incubated with a test peptide from the

30 peptide, the alanine substitution having disrupted the

binding interaction.

groups listed above at concentrations ranging from $10\,\mu\text{M}$ to $0.0045\,\mu\text{M}$. If the test peptide competes with the immobilized immunogen for binding to the antibody, the ELISA will reveal little or no binding of the antibody to the plate.

The epitope recognized by gp41-directed antibodies can be mapped in a similar manner using an alanine substitution series based on the sequence of all or part of gp41. It is also possible to use peptides whose sequence corresponds to a portion of gp41.

V3 Loop-Directed Antibodies

Described below are two antibodies which recognizes sequences within the V3 loop of HIV-MN gp120 and which can be used to generate heteroconjugate

15 antibodies.

Hybridomas F59 and F83 were generated from immunization of BALB/C mice with the closed loop immunogen RP70 (Table 1). Antibodies, designated F59/P5B3 (59.1), and F83/P6F12 (83.1) were identified as 20 antibodies which are not strain specific. Alaninesubstituted peptides that were capable of competing with RP70 for binding to the 59.1 antibody did not contain alanine substitutions within the G-P-G-R-A-F sequence. Similarly, peptides that contained the G-P-G-R-A-F 25 sequence were able to compete with RP70 for binding to the 59.1 antibody, while those that did not contain this sequence (i.e., RP129 and RP175) were not able to compete. These results indicate that the 59.1 antibody recognizes the G-P-G-R-A-F epitope. This sequence is 30 present in a wide range of HIV variants. The strain specificity of the 59.1 antibody was analyzed using the above-described techniques. These assays indicated that 59.1 recognizes the V3 loop of HIV-MN, HIV-SF2, HIV-WMJ2 and HIV-III.

ELISA assays demonstrated that the 83.1 antibody recognizes the I-X-I-G-P-G-R epitope (where X is any amino acid). The strain specificity of the 83.1 antibody was analyzed using the above-described techniques. These assays indicated that 83.1 recognizes the V3 loop of HIV-MN, HIV-Alabama, HIV-SF2, HIV-WMJ2, and HIV-Duke 7887-7.

Thus, we have identified and characterized two antibodies which recognize a number of HIV strains. One 83.1 recognizes the epitope I-X-I-G-P-G-R (where X is any 10 amino acid. Another, 59.1, recognizes the epitope GPGRAF. THe 59.1 antibody was used to generate a heteroconjugate antibody using the method of Scott et al. (J. Immunology 140:8, 1988).

gp41-Directed Antibodies

15 Recombinant gp160 was used to generate monoclonal antibodies essentially as described above. ELISA assays demonstrated that one of these antibodies, 6C5, recognizes the portion of gp41 from amino acids 584 to 611. This portion of gp41 does not vary significantly 20 from one HIV strain to another. Accordingly antibodies directed against this region are not expected to be strain specific. This antibody was used to generate and purify heteroconjugate antibodies using the method of Scott et al. (J. Immunology 140:8, 1988).

25 OKT3/59.1 and OKT3/6C5 Heteroconjugate Antibodies

The experiments described below illustrate the effect of two heteroconjugate antibodies, OKT3/59.1 and OKT3/6C5, on viral replication in CEM-ss cells (American Type Culture Collection, Bethesda, MD: Accession No. CCL119) infected with various strains of HIV. OKT3/59.1 heteroconjugate antibody was produced by covalently cross-linking an anti-CD3 monoclonal antibody, OKT3, to a second monoclonal antibody, 59.1, directed against an epitope within the V3 domain of the gp120 subunit of HIV-

OKT3/6C5 heteroconjugate antibody was produced by covalently cross-linking OKT3, to a monoclonal antibody, 6C5, directed against a conserved epitope within residues 584-611 of the gp41 subunit of HIV (numbering according 5 to Ratner et al., Nature 313:277, 1985). CD3 is a receptor closely associated with the T cell receptor for antigen (TCR). When the infected cells are grown in the presence of cytotoxic T-lymphocytes, which express the CD3 receptor, these heteroconjugate antibodies of the 10 invention dramatically decrease viral replication as measured by viral reverse transcriptase activity. Because reverse transcriptase activity is a sensitive measure of HIV activity, these results indicate that the number of virally infected cells is sharply decreased. 15 Without being bound to a particular theory, it appears that the heteroconjugate antibodies are promoting killing of infected cells by linking T lymphocytes to infected cells.

Generation of CTL

The CTL line (1F8) used for testing the activity 20 of heteroconjugate antibodies was prepared by a modification of the method of Scott et al. (J. Immunology 140:8, 1988). Briefly, donor PBL were incubated in bulk culture with an allogenic EBV-transformed lymphoblastoid 25 cell line (stimulator cells) for 7 days in RPMI 1640 medium supplemented with 20% FBS (Gibco/BRL, Grand Island, NY), supernatant derived from PHA-stimulated PBL, and 100 U/ml of recombinant interleukin-2. The cells were then cloned by limiting dilution (1 cell/well) 30 in U-bottom trays. Irradiated autologous PBL and stimulator cells were used as feeders. The clones were screened for CTL activity (assessed by lysis of stimulator cells) and NK activity (lysis of K562 cells; CCL 243, American Type Culture Collection, Bethesda, MD).

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Clone 1F8 possessing CTL activity and not NK activity was selected.

OKT3/59.1 Decreases Viral Replication as Measured by Reverse Transcriptase Activity in HIV-MN and HIV-III_B
5 Infected Cells

Unless otherwise noted, CEM-ss cells (15,000 cells/well in a 96 well plate) were infected with 64 infectious units (IU) of HIV-III_B or HIV-MN. At 18 hr post-infection, effector cells (1F8 cells 45,000/well)

10 along with varying concentrations of either OKT3/59.1 heteroconjugate or an equivalent amount of the unconjugated antibodies were added to the infected CEM-ss cells. The CTL were grown in RPMI 1640 with 10% FBS; Gibco/BRL), and were washed with fresh medium prior to addition of antibodies. After 7 days cell-free culture supernatants were harvested and assayed for reverse transcriptase activity by the method of Willey et al. (J.

Referring to Figs. 1 and 2, OKT3/59.1

20 heteroconjugate (filled diamonds) at 0.5 ng/ml
essentially eliminated reverse transcriptase activity in
CEM-ss cells infected with either HIV-III_B or HIV-MN. A
mixture of unconjugated OKT3 antibody and 59.1 antibody
(open squares) had no effect on the reverse transcriptase
25 activity even at 2,000 ng/ml.

Virol. 62:139, 1988).

Referring to Fig. 3, a separate experiment demonstrated that OKT3/59.1 heteroconjugate has no effect on reverse transcriptase activity in the absence of CTL cells (filled circles). Similarly, CTL in absence of OKT3/59.1 heteroconjugate (filled triangle) have no substantial effect on the reverse transcriptase activity of HIV-III_B infected cells. Unconjugated antibodies in the absence of CTL (open circles) and CTL alone (open triangle) have no substantial effect on the reverse transcriptase activity of infected cells. Uninfected

cells (filled square) have no detectable reverse transcriptase activity. In all cases CTL and/or antibodies were added 18 hr post-infection.

The period of HIV infection prior to the addition 5 of antibody and CTL was varied to determine whether longer periods of viral replication and thus increased viral spread affects the efficacy of the heteroconjugate molecule. CEM-ss cells were incubated with HIV-IIIB or HIV-MN (64 IU) for 6, 18, 48 or 72 hours prior to the 10 addition of antibody and CTL. Reverse transcriptase activity was measured 7 days post-infection. When infection proceeded for 6, 18 or 48 hours prior to the addition of OKT3/59.1 heteroconjugate and CTL, 0.5 ng/ml of heteroconjugate was sufficient to completely eliminate 15 reverse transcriptase activity. Under the same conditions, a mixture of unconjugated OKT3 antibody and 59.1 antibody at more than 2,000 ng/ml was required to eliminate reverse transcriptase activity. If infection was allowed to proceed for 72 hours prior to addition of 20 antibody and CTL, 1 ng/ml OKT3/59.1 was required to abolish reverse transcriptase activity. At this time point unconjugated antibody at more than 2,000 ng/ml was required to eliminate reverse transcriptase activity.

The in vitro potency of OKT3/59.1 heteroconjugate
25 was further characterized by an experiment in which the
concentration of antibody was held constant, but the
ratio of CTL to CEM-ss cells was varied. In this
experiment the concentration of OKT3/59.1 heteroconjugate
(1 µg/ml), the number of CEM-ss cells (15,000/well) and
30 the infectious dose (64 IU) of HIV-III_B or HIV-MN were
held constant, and the number of CTL added to the
cultures was varied. CTL and/or heteroconjugate antibody
were added 18 hr post-infection and reverse transcriptase
activity was measured 7 days post infection. Referring
35 to Fig. 4, reverse transcriptase activity was completely

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eliminated at a CTL:CEM-ss ratio of 0.1:1 and was partially eliminated (≥ 60%) at CTL:CEM-ss ratios as low as .006:1. This result demonstrates that OKT3/59.1 heteroconjugate is effective even when the number of target cells (CEM-ss) is significantly larger than the number of effector cells, a condition comparable to that observed in vivo.

OKT3/59.1 is Effective Against Many HIV Strains

To test whether the OKT3/59.1 heteroconjugate is 10 effective against a variety of HIV strains, CEM-ss cells (150,000/well in 24 well plates) infected with 100-1000 IU of HIV. At the time of infection a 3-fold excess of CTL (450,000/well) and 1 μ g/ml of OKT3/59.1 heteroconjugate (or 1 μ g/ml of a mixture of monomeric 15 antibodies) were added to the culture. Cultures were split 3 times per week and culture supernatants were collected at four or five day intervals for assay of reverse transcriptase activity. After the initial addition of antibody no further antibody was added. 20 splitting the culture decreases the antibody concentration and the absolute number of target and effector cells. As a control, CEM-ss were cultured with virus only. The HIV isolates tested and their V3 sequences are listed in Table 2.

Table 2: V3 Loop Sequences

	Isolate	V3 Sequence
5	MN · ·	K R K R I H I <u>G P G R A F</u> Y T T K
	Alabama	- K S H R
	Duke 6587-5	V - N H
	SF2	T S - Y H G
10	WAJ2	V - R S L S R - R E
	III _B	K S I Q R V - I G
	DUKE 6587-7	T G I - A - G
	DUKE 7887-7	TSRG-RILA-E

15 In this table a "-" indicates that the amino acid at that position is the same as in MN. The conserved GPGRAF motif in underlined

Referring to Fig. 6, compared to CEM-ss without added antibody or CTL (filled circles), OKT3/59.1

20 (unfilled squares) inhibited reverse transcriptase activity more than 95% in all cases in which the HIV isolate has the GPGRAF sequence (MN, Alabama, Duke 6587-5, III_B, SF2, and WMJ2; panels A, B, D, F, G, and H respectively). Two isolates, Duke 6587-7 (panel E) and Duke 7887-7 (panel C) having a GPGRAI motif were tested, and one (Duke 6587-7) was inhibited. A mixture of unconjugated OKT3 and 59.1 (filled triangles) had no effect.

OKT3/6C5 Heteroconjugate Inhibits Reverse Transcriptase 30 Activity of Infected Cells

OKT3/6C5 heteroconjugate was tested for its ability to inhibit reverse transcriptase activity of HIV-IIIB infected CEM-ss. Briefly, CEM-ss (15,000/well in a 96 well plate) were exposed to 64 IU of HIV-IIIB. After 18 hrs., CTL (45,000/well) and OKT3/6C5 heteroconjugate at various concentrations were added. Reverse transcriptase activity was measured 7 days later as described above.

Referring to Fig. 6, OKT3/6C5 heteroconjugate (open circles) essentially eliminated reverse transcriptase activity at concentrations as low as 0.5 ng/ml. In contrast, a mixture unconjugated OKT3 and 6C5 had no significant effect on reverse transcriptase activity.

OKT3/59.1 and OKT3/6C5 Are Cytotoxic in the Presence of CTL

antibodies, we initially performed ⁵¹Cr release assays in a model system using recombinant vaccinia virus-infected CV1 cells. Recombinant vaccinia virus which express either the HIV-III_B env gene (VPE16) or HIV-MN (VMN) env gene were used to infect CV1 cells. A recombinant vaccinia virus which does not express an HIV env gene (VSC8) was used as a negative control. Mixed cell cultures were set up essentially as described for the reverse transcriptase assays. 1F8 cells were used as effector cells and were not by themselves cytotoxic to CV1 cells or vaccinia virus-infected CV1 cells.

Referring to Table 3, OKT3/59.1 heteroconjugate at 10 µg/ml lysed 58% of the VPE16 infected CV1 cells and 62% of the VMN infected CV1 cells. Lyses of uninfected cells was very low. Cell lyses was similarly low when monomeric antibodies were added (data not shown). That the maximum lysis using OKT3/59.1 was 60% as opposed to 100% is probably the result of incomplete infection of CV1 cells by the vaccinia virus (Syncytia formation assays with limiting dilutions of CV1 cells indicated that approximately 50-60% of the cells expressing gp160). Heteroconjugates formed using 1C1 or 7C6, two antibodies which recognize epitopes at the carboxyl-terminus ofgp120 and which bind to cells expressing HIV env (as assessed by FACS), were relatively ineffective in lysing cells infected with vaccinia virus expressing either HIV-MN or

 ${
m HIV-III_B}$ env protein. Apparently cell surface binding, although necessary, is not a sufficient characteristic by which to determine whether any given antibody can generate a cytotoxic heteroconjugate antibody.

5 Table 3: Cell Lysis by Heteroconjugate Antibodies

10	- <u>Heteroconjugate</u> OKT3/59.1	Perce CV1-VSC 6	entage Cell 8CV1-VPE1 58	Lysis CV1-VMN 62
	OKT3/7C6	9	17	N.D.
15	OKT3/1C1	1	18	N.D.

Assay of Heteroconjugate Potency

The assay described below is used to determine the potency of the heteroconjugate antibodies of the invention. By using a variety of HIV strains the breadth of reactivity of a given heteroconjugate antibody may also be determined. In order to accurately determine the potency of the heteroconjugate antibodies of the invention it is important to measure the effect of the heteroconjugate antibody on HIV infected cells under carefully controlled conditions. The preferred assay is described below.

30 CEM-ss cells (150,000 cells/well in 24 well
microtiter plates, 2 ml wells) are infected with 1001000 infectious units (IU) of the desired HIV strain. At
18 hr post-infection, 450,000 effector cells are added
along with enough heteroconjugate antibody to make the
35 desired initial heteroconjugate antibody concentration.
The cells are grown under standard conditions and are
split every 3 days. No additional heteroconjugate
antibody is added, thus the heteroconjugate antibody

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concentration is halved each time the culture is split.

A control culture is set up and grown under exactly the same conditions but without heteroconjugate antibody.

The reverse transcriptase activity of both cultures is

5 measured 10 days post infection using the method of Willey et al. (J. Virology 62:139, 1988). For HIV stains which take longer than 14 days post-infection to reach peak virus production, reverse transcriptase activity should not be measured at 10 days post-infection.

10 Instead the reverse transcriptase activity should be measured at a time which is close to that of maximum virus production.

The infectious units are determined according to the Kärber method. Because viral titer can decrease during storage, it is important that the viral stock be freshly titered. Viral stocks should be carefully prepared so that the number of defective viral particles is low. For example, the multiplicity of infection for preparing viral stocks should be 0.001, cells should be grown under conditions which allow logarithmic cell growth, and virus should be collected at the peak of virus production (as determined by maximal reverse transcriptase activity or p24 expression).

Engineered Heteroconjugate Antibodies

Since, for the most part, monoclonal antibodies are produced in species other than humans, they are often immunogenic to humans. In order to successfully use heteroconjugate antibodies in the treatment of humans, it may be necessary to create chimeric antibody molecules

wherein the antigen binding portion (the variable region) is derived from one species, and the portion involved with providing structural stability and other biological functions (the constant region) is derived from a human antibody. Methods for producing chimeric antibodies in which the variable domain is derived from one species and

the constant domain is derived from a second species are well known to those skilled in the art. See, for example, Neuberger et al., WO Publication No. 86/01533, priority September 3, 1984; Morrison et al, EP 5 Publication No. 0,173,494, priority August 27, 1984. An alternative method, in which an antibody is produced by replacing only the complementarity determining regions (CDRs) of the variable region with the CDRs from an immunoglobulin of the desired antigenic specificity, is 10 described by Winter (GB Publication No. 2,188,638, priority March 27, 1986). Murine monoclonals can be made compatible with human therapeutic use by producing an antibody containing a human Fc portion (Morrison, Science 229:1202, 1985). Single polypeptide chain antibodies are 15 also more easily produced by recombinant means than are conventional antibodies. Ladner et al. (U.S. Patent No. 4,946,778) describes methods for producing single polypeptide chain antibodies and these methods may be adapted to produce heteroconjugate antibodies. 20 Established procedures would allow construction, expression, and purification of such a hybrid monoclonal antibody. Quadromas can be used to generate bispecific antibodies (Reading et al., U.S Patent Nos. 4,474,893 and 4,714,681).

25 <u>Use</u>

The antibody of the invention is administered parenterally, either via the intravenous or intramuscular route. A typical treatment regimen would comprise administration of an effective amount of antibody

30 administered over between about one week and about 6 months. The number of treatments required to control a patient's disease may vary from individual to individual, depending upon the severity and stage of the illness and the individual characteristics of each patient being

35 treated. The total dose required for each treatment may

be administered by multiple doses or in a single dose. The human monoclonal antibody may be administered alone or in conjunction with other HIV treatments, such as AZT, in order to control a patient's disease.

Pharmaceutical compositions of heteroconjugate antibodies are produced according to the intended mode of administration and may include: liposomes, solutions, suspensions and microparticles.

In some circumstances it may be desirable to

10 administer the heteroconjugate antibody along with the
appropriate effector cell (Nitta et al., The Lancet
335:368, 1990). For example, peripheral blood
lymphocytes (PBL) may be collected from an individual in
need of treatment for HIV infection (or a compatible
donor) and incubated with a heteroconjugate antibody
prior to reinfusion of the cells. In some cases the PBL
may be expanded in culture (Rosenberg et al., Science
233:1318, 1986). The PBL may also be incubated with
interleukins, interferons, or other immunomodulators. In
20 addition the cells may be incubated with molecules such
as receptor specific antibodies which will stimulate the
cytolytic activity of the effector cells (Scott et al.,
Cellular Immunology 114:370, 1988).

What is claimed is:

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SEQUENCE LISTING

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(i) APPLICANT: Repligen Corporation

5

(ii) TITLE OF INVENTION: HETEROCONJUGATE ANTIBODIES FOR

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(iii) NUMBER OF SEQUENCES: 24

10 (iv) CORRESPONDENCE ADDRESS:

(A) ADDRESSEE: Fish & Richardson

(B) STREET: 225 Franklin Street

(C) CITY: Boston

15 (D) STATE: Massachusetts

(E) COUNTRY: U.S.A.

(F) ZIP: 02110-2804

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(A) MEDIUM TYPE: 3.5" Diskette, 1.44 Mb

(B) COMPUTER: IBM PS/2 Model 50Z or 55SX

(C) OPERATING SYSTEM: IBM P.C. DOS (Version 3.30)

(D) SOFTWARE: WordPerfect (Version 5.0)

25

(vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER:

(B) FILING DATE:

30 (C) CLASSIFICATION:

(vii) PRIOR APPLICATION DATA:

(A) APPLICATION NUMBER: 07/699,773

(B) FILING DATE: 14-May-1991

35

(viii) ATTORNEY/AGENT INFORMATION:

Paul T. Clark (A) NAME: 5 (B) REGISTRATION NUMBER: 30,162 (C) REFERENCE/DOCKET NUMBER: 00231/055W01 (ix) TELECOMMUNICATION INFORMATION: 10 (617) 542-5070 (A) TELEPHONE: (617) 542-8906 (B) TELEFAX: 200154 (C) TELEX: (2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 15 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: (B) TYPE: amino acid 20 (D) TOPOLOGY: linear (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1: 25 Gly Pro Gly Arg Ala Phe (2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 2: 30 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: (B) TYPE: amino acid (D) TOPOLOGY: linear

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(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:.

Ile Xaa Ile Gly Pro Gly Arg

5

5

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 3:

(i) SEQUENCE CHARACTERISTICS:

10 (A) LENGTH:

28

(B) TYPE:

amino acid

(D) TOPOLOGY:

linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:

15

Gln Ala Arg Ile Leu Ala Val Glu Arg Tyr Leu Lys Asp Gln Gln Leu
5 10 15

Leu Gly Ile Trp Gly Cys Ser Gly Lys Leu Ile Cys
20 25

- (2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 4:
 - (1) SEQUENCE CHARACTERISTICS:

25

(A) LENGTH:

17

(B) TYPE:

amino acid

(D) TOPOLOGY:

linear

30 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:

Lys Arg Lys Arg Ile His Ile Gly Pro Gly Arg Ala Phe Tyr Thr Thr

5

10

15

35 Lys

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: (i) SEQUENCE CHARACTERISTICS: 5 (A) LENGTH: (B) TYPE: amino acid (D) TOPOLOGY: linear 10 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5: Ile Gly Pro Gly Arg 15 (2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: (i) SEQUENCE CHARACTERISTICS: 20 (A) LENGTH: 24 (B) TYPE: amino acid (D) TOPOLOGY: linear 25 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6: Tyr Asn Lys Arg Lys Arg Ile His Ile Gly Pro Gly Arg Ala Phe Tyr 30 Thr Thr Lys Asn Ile Ile Gly Cys 20 (2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 35 (i) SEQUENCE CHARACTERISTICS:

- 42 -

(A) LENGTH: 11 amino acid (B) TYPE: linear (D) TOPOLOGY: 5 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7: Ile His Ile Gly Pro Gly Arg Ala Phe Tyr Thr 10 10 (2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 8: (i) SEQUENCE CHARACTERISTICS: 15 40 (A) LENGTH: amino acid (B) TYPE: linear (D) TOPOLOGY: (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8: 20 Ile Asn Cys Thr Arg Pro Asn Tyr Asn Lys Arg Lys Arg Ile His Ile Gly Pro Gly Arg Ala Phe Tyr Thr Thr Lys Asn Ile Ile Gly Thr Ile 30 25 25 Arg Gln Ala His Cys Asn Ile Ser 35 40 (2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 9: 30 (i) SEQUENCE CHARACTERISTICS: 21 (A) LENGTH: amino acid (B) TYPE: linear (D) TOPOLOGY: 35

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:

Ser Gly Gly Thr Arg Lys Gly Ile His Ile Gly Pro Gly Arg Ala Ile 5 10 Tyr Gly Gly Ser Cys

20

- (2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 10:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH:

20

(B) TYPE:

amino acid

15 (D) TOPOLOGY:

- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:
- 20 Ser Gly Gly Thr Arg Lys Ser Ile Ser Ile Gly Pro Gly Arg Ala Phe 15 Gly Gly Ser Cys 20

25

10

- (2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 11:
 - (i) SEQUENCE CHARACTERISTICS:

30 (A) LENGTH: 19

(B) TYPE:

amino acid

(D) TOPOLOGY:

linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:

35

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Ser Gly Gly His Ile Gly Pro Gly Arg Ala Phe Tyr Ala Thr Gly Gly 15 10 Gly Ser Cys 5 (2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: (i) SEQUENCE CHARACTERISTICS: 10 (A) LENGTH: 10 amino acid (B) TYPE: (D) TOPOLOGY: linear (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12: 15 Cys His Ile Gly Pro Gly Arg Ala Phe Cys 10 5 20 (2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 13: (i) SEQUENCE CHARACTERISTICS: 25 25 (A) LENGTH: amino acid (B) TYPE: (D) TOPOLOGY: linear (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 13: 30 Asn Asn Thr Arg Lys Ser Ile Arg Ile Gln Arg Gly Pro Gly Arg Ala 15 10 5 Phe Val Thr Ile Gly Lys Ile Gly Cys 20 35

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 14: (i) SEQUENCE CHARACTERISTICS: 5 26 (A) LENGTH: amino acid (B) TYPE: (D) TOPOLOGY: linear (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 14: 10 Cys Asn Asn Thr Arg Lys Ser Ile Arg Ile Gln Arg Gly Pro Gly Arg 15 Ala Phe Val Thr Ile Gly Lys Ile Gly Cys 20 25 (2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: (i) SEQUENCE CHARACTERISTICS: 20 11 (A) LENGTH: amino acid (B) TYPE: (D) TOPOLOGY: linear 25 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 15: Ile Thr Lys Gly Pro Gly Arg Val Ile Tyr Cys 30 (2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: (i) SEQUENCE CHARACTERISTICS: 35

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17:

- 46 -16 (A) LENGTH: amino acid (B) TYPE: linear (D) TOPOLOGY: (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 16: Cys Lys Arg Ile His Ile Gly Pro Gly Arg Ala Phe Tyr Thr Thr Cys 10 5 (2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER:

(i) SEQUENCE CHARACTERISTICS:

17 (A) LENGTH: 15

> amino acid (B) TYPE:

linear (D) TOPOLOGY:

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 17:

20

5

10

Lys Arg Lys Arg Ile His Ile Gly Pro Gly Arg Ala Phe Tyr Thr Thr 10 15 5

Lys

25

- (2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 18:
 - (i) SEQUENCE CHARACTERISTICS:

17 (A) LENGTH: 30

> amino acid (B) TYPE:

(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 18:

35

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Arg 5 (2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 19: (i) SEQUENCE CHARACTERISTICS: (i) SEQUENCE CHARACTERISTICS: (i) SEQUENCE CHARACTERISTICS: (ii) SEQUENCE DESCRIPTION: SEQ ID NO: 19: Val Arg Asn Arg Ile His Ile Gly Pro Gly Arg Ala Phe His Thr Thr 5 10 15 20 Lys (2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 20: (i) SEQUENCE CHARACTERISTICS: 25 (ii) SEQUENCE CHARACTERISTICS: 25 (iii) SEQUENCE CHARACTERISTICS: 25 (iv) TYPE: amino acid (iv) TOPOLOGY: linear 30 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20: Thr Arg Lys Ser Ile Tyr Ile Gly Pro Gly Arg Ala Phe His Thr Thr 5 10 15		Lys	Lys Ser Arg	g Ile His	Ile Gly	Pro Gly A	rg Ala P	he His	Thr	Thr
(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 19: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 17 (B) TYPE: amino acid (D) TOPOLOGY: linear 15 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 19: Val Arg Asn Arg Ile His Ile Gly Pro Gly Arg Ala Phe His Thr Thr 5 10 15 20 Lys (2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 20: (i) SEQUENCE CHARACTERISTICS: 25 (A) LENGTH: 17 (B) TYPE: amino acid (D) TOPOLOGY: linear 30 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20:				5		10			15	
(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 19: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 17 (B) TYPE: amino acid (D) TOPOLOGY: linear 15 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 19: Val Arg Asn Arg Ile His Ile Gly Pro Gly Arg Ala Phe His Thr Thr 5 10 15 20 Lys (2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 20: (i) SEQUENCE CHARACTERISTICS: 25 (A) LENGTH: 17 (B) TYPE: amino acid (D) TOPOLOGY: linear 30 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20:		Arg								
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 17 (B) TYPE: amino acid (D) TOPOLOGY: linear 15 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 19: Val Arg Asn Arg Ile His Ile Gly Pro Gly Arg Ala Phe His Thr Thr 5 10 15 20 Lys (2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 20: (i) SEQUENCE CHARACTERISTICS: 25 (A) LENGTH: 17 (B) TYPE: amino acid (D) TOPOLOGY: linear 30 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20:	5				·					
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 17 (B) TYPE: amino acid (D) TOPOLOGY: linear 15 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 19: Val Arg Asn Arg Ile His Ile Gly Pro Gly Arg Ala Phe His Thr Thr 5 10 15 20 Lys (2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 20: (i) SEQUENCE CHARACTERISTICS: 25 (A) LENGTH: 17 (B) TYPE: amino acid (D) TOPOLOGY: linear 30 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20:										
(A) LENGTH: 17 (B) TYPE: amino acid (D) TOPOLOGY: linear 15 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 19: Val Arg Asn Arg Ile His Ile Gly Pro Gly Arg Ala Phe His Thr Thr 5 10 15 20 Lys (2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 20: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 17 (B) TYPE: amino acid (D) TOPOLOGY: linear 30 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20:		(2)	INFORMATION	1 FOR SEQ	UENCE ID	entificatio	NUMBE	R: 19	9:	
(A) LENGTH: 17 (B) TYPE: amino acid (D) TOPOLOGY: linear 15 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 19: Val Arg Asn Arg Ile His Ile Gly Pro Gly Arg Ala Phe His Thr Thr 5 10 15 20 Lys (2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 20: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 17 (B) TYPE: amino acid (D) TOPOLOGY: linear 30 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20:										
(A) LENGTH: 17 (B) TYPE: amino acid (D) TOPOLOGY: linear 15 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 19: Val Arg Asn Arg Ile His Ile Gly Pro Gly Arg Ala Phe His Thr Thr 5 10 15 20 Lys (2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 20: (i) SEQUENCE CHARACTERISTICS: 25 (A) LENGTH: 17 (B) TYPE: amino acid (D) TOPOLOGY: linear 30 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20:			(i) SEQUEN	ICE CHARA	CTERISTI	CS:				
(B) TYPE: amino acid (D) TOPOLOGY: linear 15 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 19: Val Arg Asn Arg Ile His Ile Gly Pro Gly Arg Ala Phe His Thr Thr 5 10 15 20 Lys (2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 20: (i) SEQUENCE CHARACTERISTICS: 25 (A) LENGTH: 17 (B) TYPE: amino acid (D) TOPOLOGY: linear 30 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20:	10		4							
(D) TOPOLOGY: linear (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 19: Val Arg Asn Arg Ile His Ile Gly Pro Gly Arg Ala Phe His Thr Thr 5 10 15 20 Lys (2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 20: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 17 (B) TYPE: amino acid (D) TOPOLOGY: linear 30 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20:			Ī							
Val Arg Asn Arg Ile His Ile Gly Pro Gly Arg Ala Phe His Thr Thr 5 10 15 20 Lys (2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 20: (i) SEQUENCE CHARACTERISTICS: 25 (A) LENGTH: 17 (B) TYPE: amino acid (D) TOPOLOGY: linear 30 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20:			, ,				cid			
Val Arg Asn Arg Ile His Ile Gly Pro Gly Arg Ala Phe His Thr Thr 5 10 15 20 Lys (2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 20: (i) SEQUENCE CHARACTERISTICS: 25 (A) LENGTH: 17 (B) TYPE: amino acid (D) TOPOLOGY: linear 30 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20:			(D) I	OPOLOGY:		linear				
Val Arg Asn Arg Ile His Ile Gly Pro Gly Arg Ala Phe His Thr Thr 5 10 15 20 Lys (2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 20: (i) SEQUENCE CHARACTERISTICS: 25 (A) LENGTH: 17 (B) TYPE: amino acid (D) TOPOLOGY: linear 30 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20:	15		(mi) CROTTE	NGP DEG	TDETON	CRO ID NO	704			
20 Lys (2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 20: (i) SEQUENCE CHARACTERISTICS: 25 (A) LENGTH: 17 (B) TYPE: amino acid (D) TOPOLOGY: linear 30 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20:	13		(XI) SEGOE	MCB DESC!	AIPIION:	SEC ID NO	19:			
20 Lys (2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 20: (i) SEQUENCE CHARACTERISTICS: 25 (A) LENGTH: 17 (B) TYPE: amino acid (D) TOPOLOGY: linear 30 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20:										
20 Lys (2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 20: (i) SEQUENCE CHARACTERISTICS: 25 (A) LENGTH: 17 (B) TYPE: amino acid (D) TOPOLOGY: linear 30 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20:		Val	Arg Asn Arg	Ile His	Ile Gly	Pro Gly Ar	a Ala Pi	ne His 1	Chr '	Thr
(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 20: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 17 (B) TYPE: amino acid (D) TOPOLOGY: linear 30 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20:										
(i) SEQUENCE CHARACTERISTICS: 25 (A) LENGTH: 17 (B) TYPE: amino acid (D) TOPOLOGY: linear 30 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20: Thr Arg Lys Ser Ile Tyr Ile Gly Pro Gly Arg Ala Phe His Thr Thr				5		_			15	
(i) SEQUENCE CHARACTERISTICS: 25 (A) LENGTH: 17 (B) TYPE: amino acid (D) TOPOLOGY: linear 30 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20: Thr Arg Lys Ser Ile Tyr Ile Gly Pro Gly Arg Ala Phe His Thr Thr	20	Lys		5		_			15	
(A) LENGTH: 17 (B) TYPE: amino acid (D) TOPOLOGY: linear 30 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20: Thr Arg Lys Ser Ile Tyr Ile Gly Pro Gly Arg Ala Phe His Thr Thr	20	Lys		5	_	_			15	
(A) LENGTH: 17 (B) TYPE: amino acid (D) TOPOLOGY: linear 30 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20: Thr Arg Lys Ser Ile Tyr Ile Gly Pro Gly Arg Ala Phe His Thr Thr	20	_	INFORMATION		- JENCE IDI	10	n number	R: 20		
(A) LENGTH: 17 (B) TYPE: amino acid (D) TOPOLOGY: linear 30 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20: Thr Arg Lys Ser Ile Tyr Ile Gly Pro Gly Arg Ala Phe His Thr Thr	20	_	INFORMATION		JENCE IDI	10	n number	R: 20		
(B) TYPE: amino acid (D) TOPOLOGY: linear 30 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20: Thr Arg Lys Ser Ile Tyr Ile Gly Pro Gly Arg Ala Phe His Thr Thr	20	_		FOR SEQU		10 ENTIFICATIO	N NUMBER	R: 20		
(D) TOPOLOGY: linear (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20: Thr Arg Lys Ser Ile Tyr Ile Gly Pro Gly Arg Ala Phe His Thr Thr		_		FOR SEQU		10 ENTIFICATIO	N NUMBER	R: 20		
30 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20: Thr Arg Lys Ser Ile Tyr Ile Gly Pro Gly Arg Ala Phe His Thr Thr		_	(i) SEQUEN	FOR SEQU		10 ENTIFICATIO	n number	R: 20		
Thr Arg Lys Ser Ile Tyr Ile Gly Pro Gly Arg Ala Phe His Thr		_	(i) SEQUEN	FOR SEQUECE CHARAC		10 ENTIFICATION ES:		R: 20		
Thr Arg Lys Ser Ile Tyr Ile Gly Pro Gly Arg Ala Phe His Thr		_	(i) SEQUEN (A) L (B) T	FOR SEQUENCE CHARACTERS ENGTH: YPE:		10 ENTIFICATIO CS: 17 amino a		R: 20		
Thr Arg Lys Ser Ile Tyr Ile Gly Pro Gly Arg Ala Phe His Thr	25	_	(i) SEQUEN (A) L (B) T (D) T	FOR SEQUENCE CHARACE ENGTH: TPE: OPOLOGY:	TERISTIC	10 ENTIFICATIO CS: 17 amino a linear	cid	R: 20		
	25	_	(i) SEQUEN (A) L (B) T (D) T	FOR SEQUENCE CHARACE ENGTH: TPE: OPOLOGY:	TERISTIC	10 ENTIFICATIO CS: 17 amino a linear	cid	R: 20		
	25	(2)	(i) SEQUEN (A) L (B) T (D) T	FOR SEQUENCE CHARACTERS ENGTH: TPE: OPOLOGY:	TERISTIC	10 ENTIFICATIO CS: 17 amino a linear	cid	R: 20		
15	25	(2)	(i) SEQUEN (A) L (B) T (D) T (xi) SEQUE	FOR SEQUENCE CHARACTERS ENGTH: TYPE: TOPOLOGY: NCE DESCR	TERISTIC	10 ENTIFICATIO CS: 17 amino a linear SEQ ID NO:	cid 20:):	Ph-
35 Glý	25	(2)	(i) SEQUEN (A) L (B) T (D) T (xi) SEQUE	FOR SEQUENCE CHARACTERS ENGTH: TYPE: OPOLOGY: NCE DESCR	TERISTIC	10 ENTIFICATIO CS: 17 amino a linear SEQ ID NO:	cid 20:): Thr 1	Chr

	(2) INFORMATION FOR SEQUENCE	IDENTIFICATION NUMBER:	21:
	(i) SEQUENCE CHARACTERIS	STICS:	
5	(A) LENGTH:	17	
	(B) TYPE:	amino acid	
	(D) TOPOLOGY:	linear	
10	(xi) SEQUENCE DESCRIPTION	ON: SEQ ID NO: 21:	
	Val Arg Arg Ser Lys Ser Ile G	Sly Pro Gly Arg Ala Phe	Arg Thr Arg
	5	10	15
15	Glu		
	(2) INFORMATION FOR SEQUENCE	IDENTIFICATION NUMBER:	22:
20	(i) SEQUENCE CHARACTERIS	STICS:	
	(A) LENGTH:	17	
	(B) TYPE:	amino acid	
_	(D) TOPOLOGY:	linear	
25	(xi) SEQUENCE DESCRIPTION	ON: SEQ ID NO: 22:	
	Lys Ser Ile Arg Ile Gln Arg G		
30	5 Gly	10	15
35	(2) INFORMATION FOR SEQUENCE	IDENTIFICATION NUMBER:	23:

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(i)	Sequence	CHARACTERISTICS:
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(A) LENGTH:

17

(B) TYPE:

amino acid

5

(D) TOPOLOGY:

linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 23:

10 Thr Arg Lys Gly Ile His Ile Gly Pro Gly Arg Ala Ile Tyr Ala Thr

5

10

15

Gly

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 24:

15

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH:

17

(B) TYPE:

amino acid

20

(D) TOPOLOGY:

linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 24:

25 Thr Ser Arg Gly Ile Arg Ile Gly Pro Gly Arg Ala Ile Leu Ala Thr

5

10

15

Glu

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Claims

A heteroconjugate antibody comprising a first 1 and a second portion joined together covalently, said 2 first portion comprising an antibody directed against an 3 antigen present on the surface of an effector cell of the 4 peripheral blood, said second antibody portion comprising 5 an antibody directed against a V3 loop sequence of the 6 gp120 envelope protein of HIV MN or a HIV MN viral 7 variant expressed on the surface of HIV-infected cells, 8 wherein said heteroconjugate antibody at an initial 9 concentration of 20 ng/ml in a first mixed cell culture 10 comprising said effector cells and CEM-ss cells infected 11 with HIV-MN decreases the reverse transcriptase activity 12 of said first mixed culture cell by at least 80% compared 13 to the reverse transcriptase activity of an otherwise 14 identical second mixed cell culture comprising said 15 effector cells and said CEM-ss cells infected with HIV-16 MN, wherein said effector cells are in 3-fold excess over 17 said CEM-ss cells in said first and second mixed cell 18 cultures, said reverse transcriptase activity is measured 19 ten days after infection, said heteroconjugate antibody 20 and said effector cells are added to said CEM-ss cells in 21 said first mixed cell culture 18 hours after infection, 22 and said first and second cell cultures are infected with 23 100-1000 infectious units of HIV-MN. 24

- 2. The heteroconjugate antibody of claim 1 where in said decrease in said reverse transcriptase activity of said first cell culture is greater than 90% compared to said reverse transcriptase activity of said second mixed cell culture.
- 3. A heteroconjugate antibody comprising a first and a second portion joined together covalently, said first portion comprising an antibody directed against an

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antigen present on the surface of an effector cell of the 4 peripheral blood, said second antibody portion comprising 5 an antibody directed against a V3 loop sequence of the qp120 envelope protein of HIV MN or a HIV MN viral 7 variant expressed on the surface of HIV-infected cells, 8 wherein said heteroconjugate antibody at an initial 9 concentration of 10 ng/ml in a first mixed cell culture 10 comprising said effector cells and CEM-ss cells infected 11 with HIV-MN decreases the reverse transcriptase activity 12 of said first mixed culture cell by at least 80% compared 13 to the reverse transcriptase activity of an otherwise 14 identical second mixed cell culture comprising said 15 effector cells and said CEM-ss cells infected with HIV-16 MN, wherein said effector cells are in 3-fold excess over 17 said CEM-ss cells in said first and second mixed cell 18 19 cultures, said reverse transcriptase activity is measured 20 ten days after infection, said heteroconjugate antibody 21 and said effector cells are added to said CEM-ss cells in said first mixed cell culture 18 hours after infection, 22 and said first and second cell cultures are infected with 23 100-1000 infectious units of HIV-MN. 24

4. A heteroconjugate antibody comprising a first 1 and a second portion joined together covalently, said 2 first portion comprising an antibody directed against an 3 antigen present on the surface of an effector cell of the 4 5 peripheral blood, said second antibody portion comprising an antibody directed against a V3 loop sequence of the 6 7 gp120 envelope protein of HIV MN or a HIV MN viral variant expressed on the surface of HIV-infected cells, 8 wherein said heteroconjugate antibody at an initial 9 concentration of 5 ng/ml in a first mixed cell culture 10 comprising said effector cells and CEM-ss cells infected 11 with HIV-MN decreases the reverse transcriptase activity 12. of said first mixed culture cell by at least 80% compared 13

24

- to the reverse transcriptase activity of an otherwise 14 identical second mixed cell culture comprising said 15 effector cells and said CEM-ss cells infected with HIV-16 MN, wherein said effector cells are in 3-fold excess over 17 said CEM-ss cells in said first and second mixed cell 18 cultures, said reverse transcriptase activity is measured 19 ten days after infection, said heteroconjugate antibody 20 and said effector cells are added to said CEM-ss cells in 21 said first mixed cell culture 18 hours after infection, 22 and said first and second cell cultures are infected with 23 100-1000 infectious units of HIV-MN.
- A heteroconjugate antibody comprising a first 5. 1 and a second portion joined together covalently, said 2 first portion comprising an antibody directed against an 3 antigen present on the surface of an effector cell of the 4 peripheral blood, said second antibody portion comprising 5 an antibody directed against a V3 loop sequence of the 6 gp120 envelope protein of HIV MN or a HIV MN viral 7 variant expressed on the surface of HIV-infected cells, 8 wherein said heteroconjugate antibody at an initial 9 concentration of 1 ng/ml in a first mixed cell culture 10 comprising said effector cells and CEM-ss cells infected 11 with HIV-MN decreases the reverse transcriptase activity 12 of said first mixed culture cell by at least 80% compared 13 to the reverse transcriptase activity of an otherwise 14 identical second mixed cell culture comprising said 15 effector cells and said CEM-ss cells infected with HIV-16 MN, wherein said effector cells are in 3-fold excess over 17 said CEM-ss cells in said first and second mixed cell 18 cultures, said reverse transcriptase activity is measured 19 ten days after infection, said heteroconjugate antibody 20 and said effector cells are added to said CEM-ss cells in 21 said first mixed cell culture 18 hours after infection, 22

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- 23 and said first and second cell cultures are infected with
- 24 100-1000 infectious units of HIV-MN.
- 1 6. A heteroconjugate antibody comprising a first
- 2 and a second portion joined together covalently, said
- 3 first portion comprising an antibody directed against an
- 4 antigen present on the surface of an effector cell of the
- 5 peripheral blood, said second antibody portion comprising
- 6 an antibody directed against the amino acid sequence
- 7 GPGRAF.
- 7. A heteroconjugate antibody comprising a first
- 2 and a second portion joined together covalently, said
- 3 first portion comprising an antibody directed against an
- 4 antigen present on the surface of an effector cell of the
- 5 peripheral blood, said second antibody portion comprising
- 6 an antibody directed against the amino acid sequence
- 7 IXIGPGR, wherein X = any amino acid.
- 1 8. The heteroconjugate antibody of claim 6 or
- 2 claim 7 wherein said heteroconjugate antibody at an
- 3 initial concentration of 20 ng/ml in a first mixed cell
- 4 culture comprising said effector cells and CEM-ss cells
- 5 infected with HIV-MN decreases the reverse transcriptase
- 6 activity of said first mixed culture cell by at least 80%
- 7 compared to the reverse transcriptase activity of an
- 8 otherwise identical second mixed cell culture comprising
- 9 said effector cells and said CEM-ss cells infected with
- 10 HIV-MN, wherein said effector cells are in 3-fold excess
- 11 over said CEM-ss cells in said first and second mixed
- 12 cell cultures, said reverse transcriptase activity is
- 13 measured ten days after infection, said heteroconjugate
- 14 antibody and said effector cells are added to said CEM-
- 15 ss cells in said first mixed cell culture 18 hours after

- 16 infection, and said first and second cell cultures are
- 17 infected with 100-1000 infectious units of HIV-MN.
- 1 9. The heteroconjugate antibody of claim 1
- 2 wherein said heteroconjugate antibody at an initial
- 3 concentration of 20 ng/ml in a first mixed cell culture
- 4 comprising said effector cells and CEM-ss cells infected
- 5 with HIV-III_B decreases the reverse transcriptase
- 6 activity of said first mixed culture cell by at least 80%
- 7 compared to the reverse transcriptase activity of an
- 8 otherwise identical second mixed cell culture comprising
- 9 said effector cells and said CEM-ss cells infected with
- 10 HIV-III_B, wherein said effector cells are in 3-fold
- 11 excess over said CEM-ss cells in said first and second
- 12 mixed cell cultures, said reverse transcriptase activity
- 13 is measured ten days after infection, said
- 14 heteroconjugate antibody and said effector cells are
- 15 added to said CEM-ss cells in said first mixed cell
- 16 culture 18 hours after infection, and said first and
- 17 second cell cultures are infected with 100-1000
- 18 infectious units of HIV-III_B.
- 1 10. The heteroconjugate antibody of claim 1
- 2 wherein said heteroconjugate antibody at an initial
- 3 concentration of 20 ng/ml in three or more mixed cell
- 4 cultures each of which comprises said effector cells and
- 5 CEM-ss cells infected with one of the HIV strains:
- 6 Alabama, Duke 6587-5, Duke 6587-7, Duke 7887-7, SF2,
- 7 WMJ2, and IIIB, decreases the reverse transcriptase
- 8 activity of each of said mixed cell culture by 80%
- 9 compared to the reverse transcriptase activity of an
- 10 otherwise identical mixed cell culture comprising said
- 11 effector cells and CEM-ss cells infected with said same
- 12 strain of HIV, wherein said effector cells are in 3-fold
- 13 excess over said CEM-ss cells in said first and second

PCT/US92/03616

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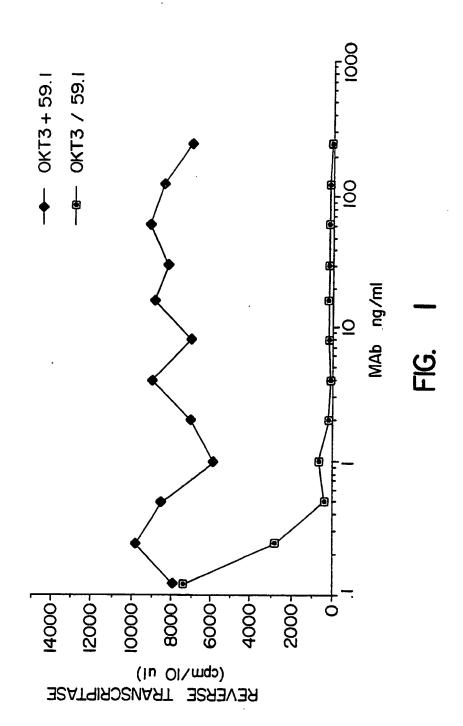
- 14 mixed cell cultures, said reverse transcriptase activity
- 15 is measured ten days after infection, said
- 16 heteroconjugate antibody and said effector cells are
- 17 added to said CEM-ss cells in said first mixed cell
- 18 culture 18 hours after infection, and said first and
- 19 second cell cultures are infected with 100-1000
- 20 infectious units of said strain of HIV.
 - 1 11. A heteroconjugate antibody comprising a
 - 2 first and a second portion joined together covalently,
 - 3 said first portion comprising an antibody directed
- 4 against an antigen present on the surface of an effector
- 5 cell of the peripheral blood, said second antibody
- 6 portion comprising an antibody directed against the amino
- 7 acid sequence QARILAVERYLKDQQLLGIWGCSGKLIC.
- 1 12. The heteroconjugate antibody of claim 11
- 2 wherein said heteroconjugate antibody at an initial
- 3 concentration of 20 ng/ml in a first mixed cell culture
- 4 comprising said effector cells and CEM-ss cells infected
- 5 with HIV-MN decreases the reverse transcriptase activity
- 6 of said first mixed culture cell by at least 80% compared
- 7 to the reverse transcriptase activity of an otherwise
- 8 identical second mixed cell culture comprising said
- 9 effector cells and said CEM-ss cells infected with HIV-
- 10 MN, wherein said effector cells are in 3-fold excess over
- 11 said CEM-ss cells in said first and second mixed cell
- 12 cultures, said reverse transcriptase activity is measured
- 13 ten days after infection, said heteroconjugate antibody
- 14 and said effector cells are added to said CEM-ss cells in
- 15 said first mixed cell culture 18 hours after infection,
- 16 and said first and second cell cultures are infected with
- 17 100-1000 infectious units of HIV-MN.

- 1 13. The heteroconjugate of claim 1 or claim 11
- 2 wherein said effector cell is chosen from the group
- 3 consisting of cytotoxic T lymphocytes, neutrophils,
- 4 monocytes/macrophages, and large granular lymphocytes.
- 1 14. The heteroconjugate of claim 1 or claim 11
- 2 wherein said antigen present on the surface of an
- 3 effector cell is CD3.
- 1 15. The heteroconjugate antibody of claim 1
- 2 wherein said heteroconjugate antibody at an initial
- 3 concentration of 200 ng/ml in a first mixed cell culture
- 4 comprising said effector cells and CEM-ss cells infected
- 5 with an HIV strain other than HIV-MN decreases the
- 6 reverse transcriptase activity of said first mixed
- 7 culture cell by at least 50% compared to the reverse
- 8 transcriptase activity of an otherwise identical second
- 9 mixed cell culture comprising said effector cells and
- 10 said CEM-ss cells infected with said HIV strain other
- 11 than HIV-MN, wherein said effector cells are in 3-fold
- 12 excess over said CEM-ss cells in said first and second
- 13 mixed cell cultures, said reverse transcriptase activity
- 14 is measured ten days after infection, said
- 15 heteroconjugate antibody and said effector cells are
- 16 added to said CEM-ss cells in said first mixed cell
- 17 culture 18 hours after infection, and said first and
- 18 second cell cultures are infected with 100-1000
- 19 infectious units of said HIV strain other than HIV-MN.
 - 1 16. The heteroconjugate antibody of claim 1
 - 2 wherein said heteroconjugate antibody binds to the V3
 - 3 loop of an HIV strain other than HIV-MN.

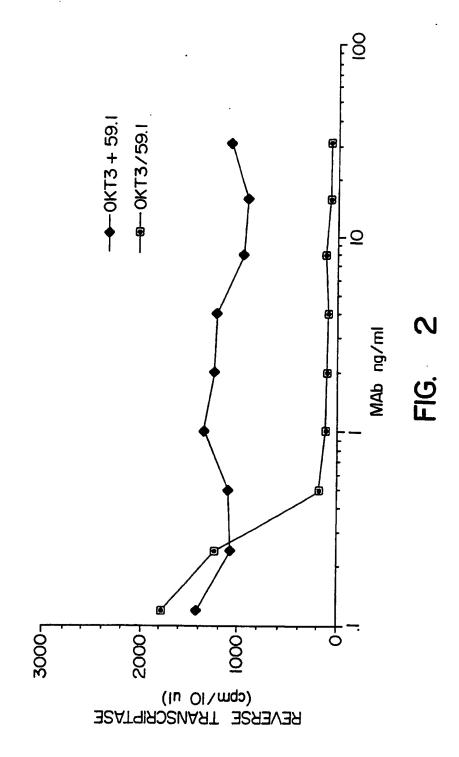
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1	17. A pharmaceutically acceptable composition
2	comprising a pharmaceutically effective amount of a
3	heteroconjugate antibody of claim 1 or claim 11.

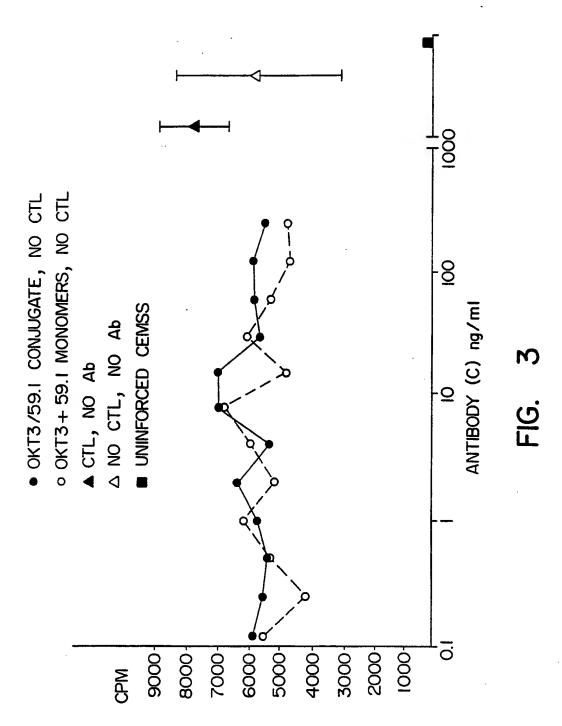
- 1 18. An HIV-targeted effector cell comprising:
- 2 (a) an effector cell expressing a cell surface
- 3 antigen; and
- 4 (b) the heteroconjugate antibody of claim 1 or
- 5 claim 11.



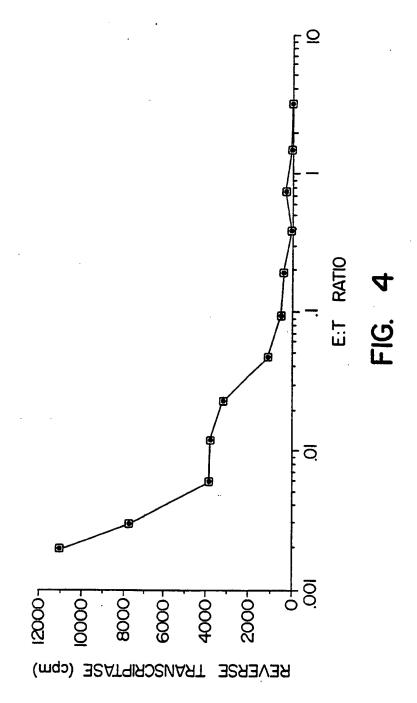
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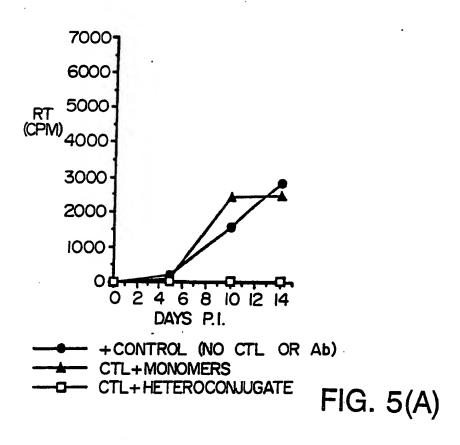


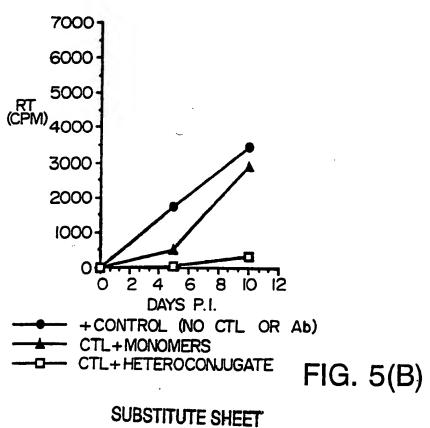
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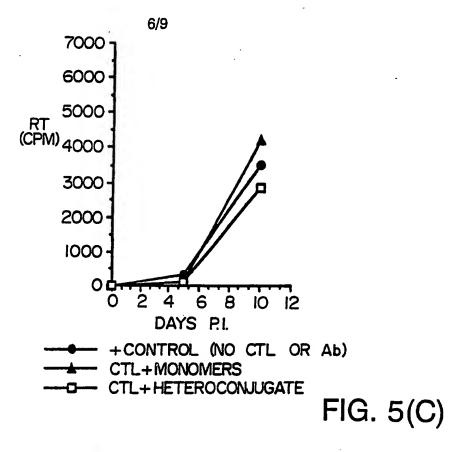


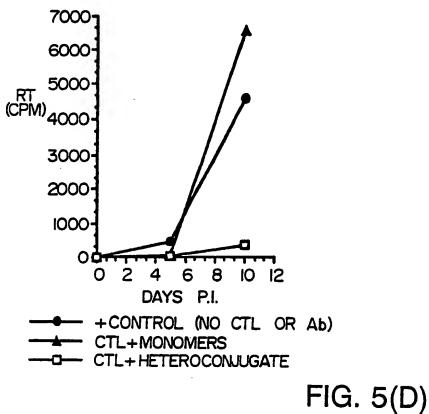
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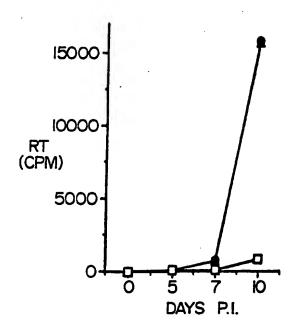








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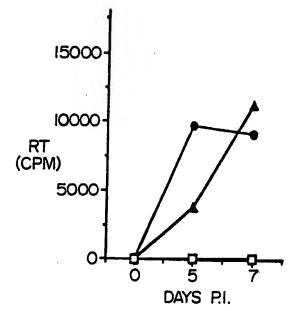


+ CONTROL (NO CTL OR Ab)

CTL+MONOMERS

CTL+ HETEROCONJUGATE

FIG. 5(E)



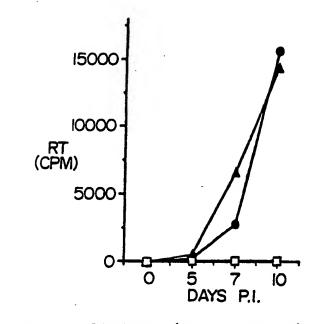
+ CONTROL (NO CTL OR Ab)

CTL+MONOMERS

CTL + HETEROCONJUGATE

FIG. 5(F)

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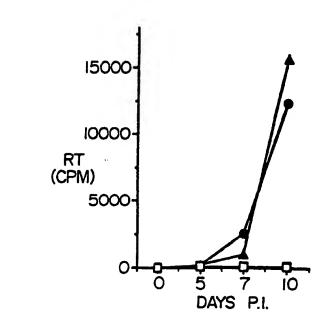


+ CONTROL (NO CTL OR Ab)

CTL+MONOMERS

CTL + HETEROCONJUGATE

FIG. 5(G)



+ CONTROL (NO CTL OR Ab)
CTL+MONOMERS

CTL + HETEROCONJUGATE

FIG. 5(H)

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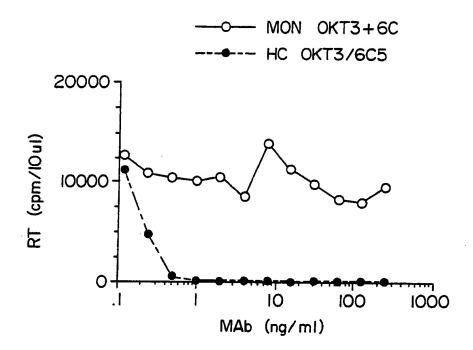


FIG. 6

INTERNATIONAL SEARCH REPORT

International application No. PCT/US92/03616

A. CLASSIFICATION OF SUBJECT MATTER IPC(5) : Please See Extra Sheet. US CL : Please See Extra Sheet. According to International Patent Classification (IPC) or to both national classification and IPC
According to International Patent Classification (IPC) or to both national classification and IPC
B. FIELDS SEARCHED
Minimum documentation searched (classification system followed by classification symbols)
U.S. : Please See Extra Sheet.
Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched
Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)
AUTOMATED PATENT SYSTEM, FILE: USPAT; DIALOG, FILE:BIOSIS, CAS, MEDLINE, PASCAL, WPI, EMBASE
SEARCH TERMS:
C. DOCUMENTS CONSIDERED TO BE RELEVANT
Category* Citation of document, with indication, where appropriate, of the relevant passages Relevant to claim No.
Y BIOTECHNOLOGY, VOLUME 3, ISSUED OCTOBER 1985, CHANG ET AL, 1-6, 8-10,
DETECTION OF ANTIBODIES TO HUMAN T-CELL LYMPHOTROPIC VIRUS-III AND 13-17
(HTLV-III) WITH AN IMMUNOASSAY EMPLOYING A RECONIANNAT
ESCHERICIA COLI-DERIVED VIRAL ANTIGENIC PEPTIDE*, PAGES 905-909, ENTIRE DOCUMENT
Y VIRILOGY, VOLUME 164, ISSUED 1988, GURGO ET AL, "ENVELOPE SEQUENCES 1-6, 8-10,
OF TWO NEW UNITED STATES HIV-1 ISOLATES*, PAGES 531-536, ENTIRE AND 13-17 DOCUMENT.
Y JOURNAL OF VIROLOGY, VOLUME 61 NO. 2, ISSUED FEBRUARY 1987, 1-6, 8-10,
MODROW ET AL, "COMPUTER ASSISTED ANALYSIS OF ENVELOPE PROTEIN AND 13-17 SEQUENCES OF SEVEN HUMAN IMMUNODEFICIENCY VIRUS ISOLATES:
PREDICTION OF ANTIGENIC EPTIOPES IN CONSERVED AND VARIABLE
REGIONS", PAGES 570-578, ENTIRE DOCUMENT.
Y WO, A, 91/00360 (FANGER ET AL), 10 JANUARY 1991, ENTIRE DOCUMENT. 1-6, 8-10,
AND 13-17
Further documents are listed in the continuation of Box C. See patent family annex.
date and not in conflict with the application but cited to understand the
to be part of particular relevance
*E" earlier document published on or after the international filing date "L" document which may throw doubts on priority chain(s) or which is "L" document which may throw doubts on priority chain(s) or which is when the document is taken alone
cited to establish the publication date of another citation or other
special reason (as specified) "Y" document of particular relevance; the claimed invention cannot be considered to investive an inventive attep when the document is combined with one or more other such documents, such combination
means being obvious to a person skilled in the art
P document published prior to the international filing date but later than "&" document member of the same patent family the priority date claimed
Date of the actual completion of the international search Date of mailing of the international search report
23 June 1992
30ctp 4
Name and mailing address of the ISA/ Commissioner of Patents and Trademarks Box PCT Washington, D.C. 20231 Authorized officer T. MICHAEL NISBET
Box PCT Washington, D.C. 20231 T. MICHAEL NISBET
Facsimile No. NOT APPLICABLE Telephone No. (703) 308-4204

INTERNATIONAL SEARCH REPORT

International application No. PCT/US92/03616

A. CLASSIFICATION OF SUBJECT MATTER: IPC (5):

A61K 39/00, 39/42, 35/14; C12Q 1/00; G01N 33/53; C12P 21/02; C12N 9/96, 5/00, 15/00; C07K 3/00, 13/00, 15/00

A. CLASSIFICATION OF SUBJECT MATTER: US CL :

424/85.8, 86; 435/7.1, 70.21, 188.5, 240.27, 965, 972, 974; 436/819, 822; 530/387.3, 387.9, 388.3, 388.35, 388.7, 388.75, 389.4, 867; 930/221; 935/15, 93, 101, 105

B. FIELDS SEARCHED
Minimum documentation searched
Classification System: U.S.

424/85.8, 86; 435/7.1, 70.21, 188.5, 240.27, 965, 972, 974; 436/819, 822; 530/387.3, 387.9, 388.3, 388.35, 388.7, 388.75, 389.4, 867; 930/221; 935/15, 93, 101, 105

BOX II. OBSERVATIONS WHERE UNITY OF INVENTION WAS LACKING This ISA found multiple inventions as follows:

I. CLAIMS 1-6,8-10, AND 13-17 CONTAIN CLAIMS TO SEPARATE ANTIBODY CONJUGATES WHICH ARE FACILITATE THE TREATMENT OF AIDS INFECTION.

II. CLAIM 7 IS DRAWN TO A SPECIAL ANTIBODY CONJUGATE WHICH BINDS A SEPARATE SEQUENCE FROM THE ONE DESCRIBED ABOVE IN GROUP I. AS SUCH THE DIFFERENT BINDING SEQUENCE REPRESENTS A SEPARATE PRODUCT.

III. CLAIMS 11-12 ALSO DEAL WITH AN ANTIBODY CONJUGATE WHICH BINDS A SEPARATE SEQUENCE FROM EITHER THE TWO ANTIBODY CONJUGATES OF GROUPS I OR II. THEREFORE, THIS TECHNICALLY REPRESENTS A MATERIALLY SEPARATE PRODUCT.

IV. CLAIM 18 CLAIMS A MATERIALLY SEPARATE COMPOSITION OF CELLS CONTAINING THE ANTIBODY CONJUGATES OF THE PREVIOUS CLAIMS. THIS IS A SEPARATE PRODUCT AND IS PROPERLY RESTRICTABLE.